SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21775	52143	A	21903	843	2055	HHRMAQNLSRYGNDKRPEESQ FRVVNVKPVTLYDVAEYAGVS YQTVSRVVNQASHVS\AKTREK \VEAAMAELNYIPNRVAQQLAG KQSLLIGVATSSLALH\APSQIV AAIKSRADQLGASVVVSMVF\R SGLVACHTAPHNLL/ARKRGPA GLIINYPLDD\QDAIAVEA\ACT NVPALFLDVSDQTPI\NSIIFSHE DGTRLGVEHL\VALGH\QQI\AL LAGPLSSVS\ARLRLAG\WHNFL TGN\QIQPIA\ER\EGDWSAMSGF QQT\LQMLNEGIVPHWRLLVAQ RSRMALG\AMRRHYPESGLRV GA\DILRVGGNTTIPED\SSCYIP AL*PPIKQDFRLLGA/QTSVDR\L LQLSQGQAVKGN\QLLPVSL\V KRKTTL\APNTQTA\SPRALPRF
		ļ. —	21221		1201	H*MQLA\RQVSRLESGQ
21776	52144	A	21904	1	1201	
21777	52145	A	21905	(46	934	
21778 21779	52146 52147	A A	21906 21907	646	1713	MLCTKWDKLGLEKNAGKNKN
						AYKWYGWPANAPDEPHIEQLV GECRVMRQLKRLISRIAPSPSSV MVVGESGTAIPVVSHAEFKGGF ADIGVHYLDWTSRTTEKSSTKS HKDDFGYLEFEGGANFSWGEM YGFFDWENFYNGRHNKPGSEQ RYTFKNTKPYLSWVDTGFSLYL HAYGTYGSANRVNFLDDM\FL AGFGFQF*PAGGW/WGSNPFFA K\RYTRSKPYYT\GDNGYV\AG WVAGYNFMLGSEKFTLTNWN EYEFDRDATYAADFLPLYDVD CQDNGNLEYDTYSQPEWKHNL FDHYLAVLYRFKDESGKEQFSG AVVKTREATPGKEIEAITRML DFSPRLKKLADCPSPRPVFEALC ICSMLDALLLLCPLDYHASCDN YSYALLDPQGFPLPHRRLWGPY IDVDVPMLHLIQHAWAQDALL SCLYSDLLYMSSFPYPAGTLTR LMGVTSYNHKNTLLTYCITLMS TIRFAYDGDEEIAMKYTDFLKG ELSMNIRPLHDRVIVKRKEVET KSAGGIVLTGSASAKSTRGEVL AVGNGRILENGEVKPLDVKIGD IVFFNDGYRGEI
21780	52148	A	21908	2060	2404	
21781	52149	В	21909	1	1377	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21782	52150	A	21910	52	1719	IGQVVDVDTCFTRIHLVFVNFD NDTGSIHVLNDTTTFSYRSYTG VNGNSTFHTSTNQRLISAQSRN GLTLHVRTHQCTVGVIVFQERD QGRTDGYHLLGGYVHVVNLV AAEQAGFAFATASYQVFYEVA FFIQVGVRLGDNVVAFFDSRQV VNFVSYNTVGHFTIRSLKEAVF VSLCVHGQRVDQTDVRTFRGF DWTDATVVSRVYVSNFEACTF TGQTAWAECGDTTFVRNLRQR VVLVHKLRQLAGTEELFHCCG SQDPQQPWGWNLRVNPLAQTT GHAHEAQRKQWHKGNEEADD PEPERAFTPGFVQFETERFRPPV GHPCEAAEDHAADDHVMEVR
						NQEQAVMQNEVRARYGQQHA SHPAHREGDDKTDGPQHRRME HDAALIHGEQPVEDLHPGRDG DNHCGDTKEGVYVRARTHRKE VVQPNDERQHGDTDGCPYQRG ITKQTFTREGCRNFGEHTENRQ DQDVNFRVAPGPNQVDVHHH VATHIVGKEMGAQIAIQGQHAT YRQNGKDL*SNRGVASGNPQC RHRAHRRATDCQTVIKQPVFFG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21783	52151	Α	21911	608	2508	YITATSCSGWMIRWCAKAAKC
						CA/CSRGPGGKLRRHKGAEETL
						CGNTRVLIAPQLPAVIIATLSGS
						RSQTGRREGPTPTTVMIQDHTG
						SSTDDECSANVQRIFNRIMLCSI
						VSRPRTIPFHDLAQSFRQFRNVK
		ŀ				CLWCAFEGAQLMRIQPAIVMV
						KTIHRERLNSGQFRCHQEARNI
						PPHRQSRFLASRGLLAELMFML
						YGIGELPEIVTLPKEIRGNVDNS
		l				RHAVFLANSGMPELRFIRTHGS
						RFTSQDCTLFNWLARIITPVLQS
						WLNDEEQQVALRLLEKDRDHH
						RVLVDITNAVLSHLDLDDLIAD
						VAREIHHFFGLASVSMVLGDHR
						KNEKFSLWCSDLSASHCACLPR
1						CMPGESVLLTQTLQTRQPTLTH
						RADDLFLWQRDPLLLLLASNG
į						CESALLIPLTFGNHTPGALLLAH
						TSSTLFSEENCQLLQHIADRIAI
						AVGNADAWRSMTDLQESLQQ
						ENHQLSEQLLSNLGIGDIIYQSQ
						AMEDLLQQVDIVAKSDSTVLIC
ŀ	j.					GETGTGKEVIARAIHQLSPRRD
						KPLVKINCAAIPASLLESELFGH
						DKGAFTGAINTHRGRFEIADGG
						TLFLDEIGDLPLELQPKLLRVLQ
						EREIERLGGSRTIPVNVRVIAAT
						NRDLWQMVEDRGVI
21784	52152	Α	21912	147	2699	MTIRAICSCCPLPDA*NVLM*IY
						MSTFQASGNGQQLQIARIVIQNI
			1			TG
21785	52153	Α	21913	432	1016	FCPSSCLAATAPAA*AGWSFLP
						SPAEQNRWRFPLSRFCQIPAPL/
						LGTLIISLSVFCSILAASAAEIPG
						SVDGIYNKSPSLI\WVKIPRQLY
						*AAREPSAPKAPPPAAWFSASA
						VLSATA\MINTNQKAIERVTAFR
	1					RNFATDPVAHQDRDHYDRQQR
						RTRHRIRFGERQRAKQTPFLPFQ
						GEDRDKRQRDNQQTDK
						• • • • • • • • • • • • • • • • • • • •

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21786	52154	A	21914		1146	MTTSMLNAKLLPSAPSAAVVV
21780	32134	A	21914	1	11140	VRVVVVGNAPTSRKVGPEEK
			1			DWSMASSGTTSTRKRFTGAEFI
	1					VHFLEQ/PGH*DCDRHSG/S/CSI
		i				LPVYDALSQSTQIRHILARHEQ
	1					GAGFIAQGMARTDGKPAVCMA
						CSGPGATNLVTAIADARLDSIPL
						ICITGQVPASMIGTDAFQEVDT
i						YGISIPITKHNYLVRHIEELPQV
			ļ			MSDAFRIAQSGRPGPVWIDIPK
	İ					DEQTAVFEIETQPAMAEKAAAP
			1			AFSEESIRDAAAMINAAKRPVL
1		İ				YLGGGVINANARVRELAEKAQ
		1				LPTTMTLMALGMLPKAHPLSL
						GMLGMHGVRSTNYILQEADLLI
	1					VLELAGVQTVDTDVNASKPAS
	,					ATRHIAFQTVTIGCDRNLADSV
1						VFPHSVDDIGKVAA
21707	60155		21915	1001	1037	MVGMASSCMPAFTRLELYSVIS
21787	52155	Α	21913	1001	1037	1
ĺ						QSTQIRH/ILARHEQGAGFIAQG
		ļ				MA/RTDGKPAVCMACSGPG/AT
1		ŀ				NLVTAIADARLDSIP/LICITGQV
						PASMIGTDA/FQEVDTYGI/STM
						TLMALGMLPKAH/ALSLGMLG
						MHGVRSTNYFCQEADLSIVLG
					ŀ	ALFDAGQIGKTSSFCPNAKI/IH
						VDIDRAE/REFPCPIPKACDLLS
						H/YGLINAVAACVDDNAIITTD
	1					VGQHQMWTAQAYPLNRPRQW
						LTSGGLGTMGFGLPAAIGAALA
						NPDRKVLCFSGDGSLMMNIQ*
						MIIAIPTGVKIFNWCSPCIRPHR
						VPFCDAVAIGFIVTFSDVIQIASI
			}			YEIPCRDPTGVASTEYSSSCRR
			į			RRNTTLCDPGSPAQRQSPQAGQ
	i					NPSCPAHRRSATGVGD
21788	52156	Α	21916	135	449	
21789	52157	A	21917	5	214	
21790	52158	Α	21918	2	337	IGIAKHAQQLLLNVMQTRAVG
]					AEVOTVSATGSAVRQALANNL
						QPFAIFFQINA/W**RAGRYPDD
						HGISPETALYGWHTSATGTAG
						MR*SVRPWLQSGQGCSGWRRQ
]	QPRWSP
21701	52150	1	21919	1	774	At VA OI
21791	52159	Α	21919		1//4	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' ' '
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21792	52160	Α	21920	2	914	RHIRTPAHLERFIVDIALNVADG
						HRAKLIVQSAGTFAQTVLRADS
						SANFRQGVGLVRQFRRFKNTPL
						VGELQPVRDVVVYRAFPLAVW
						VAARQAAVSLRFGLAFGKRLV
						NFNKLNFADLQRFLWRINAAQ
	•					LSCATTT/ADPAVLLHHWTSV*
						PAQI*WR\S*PVRQNLFTALRAG
						MLHMLGDKTTQMLFMDRQSV
						NA*DLDQFRVNAFAEIAAFIEN
			ŧ			VSETAGHARTEVNAGFTQNAD
						DTTRHIFTTVVANAFYHGDSTG
						VTHAETFTRAACGIQTTTGRAI
						QAGVADNACFMAAEGGTNRR
						TNGD
21793	52161	A	21921	204	371	PGPAGAQSAALYRWSPGQAVS
						RADWFRFADC*IYPENRRNAGS
						AYRPRLSPAYCG
21794	52162	Α	21922	2664	2970	TPDTAPAGCRSRAPDGPSLPRQ
		1	-			MPRRRG*FHTARYCRTPRAITA
						*R*TE*STCPARTGEWYAHARY
						APGTSRQTATMPPTMPRRRASS
						YSSTQPDGRRQNC
21795	52163	A	21923	318	459	
21796	52164	Α	21924	577	888	TNSSNQQLAEGNRERLIGRNTA
						GPERITAGRR*LFSCQNRPCGW
						RFNKCHIGMPVVVIWRIIVMGI
						QHDDLVNPFGMRVNRVNMQF
						AKTLGQRALLLRRNSLIA
21797	52165	Α	21925	3	711	AGAHSGPSHPALATTAARQSV
						AGNRSGISGFSGHSRNPTRMSA
						RCL*SS\ALERLMRRLNGGEIQI
		l				SDVTTTTPS\LSPQVYCSAMSRN
						LCTRATPRWQSAGHPYCRWTA
						SYCAIYSDRSIRGNYSTRQSFAS
						GRRV\PTGSWQKTESEKDCSTC
		ĺ				CANWANDRTPLPTSTSSANCST
						NVQTSVVPVLRRWIWLTSLRV
•						VLTVSLKSVCARGTSPQASCWF
						VKRAASSATSPVVITTC
21798	52166	Α	21926	244	535	TRYPDDHGISPETALYGWHTSA
		ļ -				TGTAGMR*SVRPWLQSGQGCS
		l				GWRRQQPRWSP**RSC**CSVY
						PATL*FCRLYNQTSGPGHSDLP
						AFPDSDRRKI
L	L	<u> </u>	<u> </u>	<u> </u>	<u> </u>	rii Bobildi

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21799	52167	A	21927	869	2923	PKKDLKIATSAMMEAAYSV*SR TLFW/SPVRDSQVTDTQWQGL MSQLRLQGFDTLVLQWTRYGD AFTQPEQRTLLFKRAAAAQQA GLKLIVGLNADPEFFMHQKQSS AALESYLNRLLAADLQQARLW SAAPGITPDGWYISAEIDDLNW RSEAARQPLLTWLNNAQRLISD VSAKPVYISSFFAGNMSPDGYR QLLEHVKATGVNVWVQDGSG VDKLTAEQRERYLQASADCHD GAAPHGK\VFKLIIA*RARR
21800	52168	В	21928	72	690	
21801	52169	Α	21929	1	2241	
21802	52170	A	21930	1607	1893	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPAFFQLLIK HVD/SMCHQRLSGSRKSKHNVI AVVFSYCSVSVFSASCPTHQVL LQTQYF
21803	52171	Α	21931	1	3960	
21804	52172	Α	21932	1	1281	
21805	52173	Α	21933	1	1491	
21806	52174	Α	21934	1	1254	
21807	52175	Α	21935	1	723	``
21808	52176	A	21936	740	880	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPGFFPTVN
21809	52177	Α	21937	1	1191	
21810	52178	Α	21938	1033	1248	
21811	52179	A	21939	1	1425	
21812	52180	Α	21940	1	2784	
21813	52181	Α	21941	676	1467	
21814	52182	В	21942	1	3288	

2805

NO: of pept	tide hod	1			Amino acid sequence (X=Unknown,
		in USSN	location of first		*=Stop codon, /=possible nucleotide
sequen	ice	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		1
21815 52183	3 A	21943	145	1929	VSGVIEIADGSRRRKAAALTES
21013 32103	′ ^	21743	145	1727	DYRVLVGELDDEQMAALSRLG
					NDYRPTSAYERGQRYASRLQN
					EFAGNISALADAECDNLKTCHT
					SHGSVMAETAVINHKKRKNSP
					RIVQSNDLTEAAYSLSRDQKRM
					LYLFVDQIRKSDGTLQEHDGIC
	ŀ				EIHVAKYAEIFGLTSAEASKDIR
	ŀ	:			QALKSFAGKEVVFYRPEEDAG
					DEKGYESFPWFIK/RPSRGLYSV
					HINPYLIPFFIG\LQNRFTQFRLS
					ETKEITNPYAMRLYESLCQYRK
					PDGSGIVSLKIDWIIERYQLPQS
					YQRMPDFRRRFLQGFCRFRNH
					HQTGFSPAGANQRGPLAATLSG
					PGGEGQSAVARLTGEKKNHPG
					AQYANRLSPRVGRFINAAGTTG
					FPTWKAGSERNAINDDVTYAIK
					PTCWPGLDIIPSCLALHRIETEL
					MGKFDEGKLPTDPHLMLRLAIE
	·				TVAHDYDVIVIDSAPNLGIGTIN
					VVCAADVLIVPTPAELFDYTSA
					LQFFDMLRDLLKNVDLKGNSN
					GSQSPWMEEQIRDAWGSMVLK
					NVVRETDEVGKGQIRMRTVFE
					QAIDQRSSTGAWRNALSIWEPE
					CNEISIGVSLDQDGGSNSVLRK
21816 52184	1 A	21944	1790	2947	TDFSVRCKGCRTGPFLRLIFQDR
					IVGVELIAVF*NTGPRFSSCRLS
					WRRPFDKTCRMRWHQRLSGFP
					QEQNDGKQQPILC
21817 52185	5 A	21945	1	888	
21818 52186	5 A	21946	252	1863	RKNPFILH*LFR*TLRQTKPDDS
					AGKGVKI**HTQNQRSGRSQND
					F*RRLW
21819 52187	В	21947	1	1794	

TYRQSNGITASHYSYARN*RV.VMARGNAITLPVCGRDVKFTI. EVURGOSVEKTSRVWSGLLEL KAAASLYFLEQHKPYEEIGTSF GSVLROSVEKTSRVWSGLLEL KAAASLYFLEQHKPYEEIGTSF GSVLVSVVGEGPFVEEDLRKS] PGRWDKHKMFDAIEEPKKDF DKFDANDPILKDQTQEWSGSA TFTSDGKIRLFYTDYSGKHKMF DKFDANDPILKDQTQEWSGSA TFTSDGKIRLFYTDYSGKHKMF GSLTTAQVNVSKSDDTLKING EDHKTIFDGDGKTYQNVQFID GHKYLVFEANTGTENGYQGE GERNYTSGDMFILRDPHYVFDF GHKYLVFEANTGTENGYQGE SLFNKAYYGGGTNFFRKESQK QQSAKKADAELANGALGIELI YDYTLKKVMKAADHFQTPVT EJERANVFKMG DKFDANYYGGGTNFFRKESQK QQSAKKADAELANGALGIELI YDYTLKKVMKAADHFQTPVT EJERANVFKMG NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)	
IGSLASNLSLSLKTRPAFFQLSI LSPT							
21823 52191 B 21951 50 1690	21821	52189	A	21949	761	916	IGSLASNLSLSLKTRPAFFQLSIE
21824 52192 B 21952 I 1032	21822	52190	A	21950	1	1437	
21825 52193 B 21953 I 1974	21823	52191	В	21951	50	1690	
21826	21824	52192	В	21952	1	1032	
*SFRIGSLASNLSLSLKTRPAFFI LSIE\FRRPFDKTCRLMCHQRLS GFPHEQTQRDSTCIQLLFRQRF AAASCPTHQVLLQTQYF 21827 52195 B 21955 7 1702 21828 52196 A 21956 527 665 NKRIFPSDVKVAEPDHSCV*SF IGSLAFNLSLALKTPARVFSTVI 21829 52197 A 21957 469 687 21830 52198 A 21958 2 1018 EGSDSRAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEK QVPQFDQSTIKNIESAKGLDVW DSWPLQNADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSG KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVEI HKTIFDGDRKTYQNGQQFIDEG NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQTA GVSPTVQKLKSLKFDVRGQEE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPR\IVQSNHLI 21831 52199 B 21959 7 11113	21825	52193	В	21953	ı	1974	
21828 52196	21826	52194	A	21954	878	1202	SVYNKRIFPSDVKVAEPDHSCV *SFRIGSLASNLSLSLKTRPAFFQ LSIE\FRRPFDKTCRLMCHQRLS GFPHEQTQRDSTCIQLLFRQRF AAASCPTHQVLLQTQYF
IGSLAFNLSLALKTPARVFSTVI	21827	52195	В	21955	7	1702	
21830 52198 A 21958 2 1018 EGSDSRAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEK QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSE KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQE LTTAQVNVSKSDDTLKINGVEI HKTIFDGDRKTYQNGQQFIDEC NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQTA GVSPTVQKLKSLKFDVRGQEE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPR\IVQSNHLI			A		527		NKRIFPSDVKVAEPDHSCV*SFR IGSLAFNLSLALKTPARVFSTVD
GVSHITRHDMLQIPKQQQNEK QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSE KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVEI HKTIFDGDRKTYQNGQQFIDEC NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQTA GVSPTVQKLKSLKFDVRGQEE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPR\IVQSNHLI			$\overline{}$		469		
							EGSDSRAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVED HKTIFDGDRKTYQNGQQFIDEG NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQTA GVSPTVQKLKSLKFDVRGQEES VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPR\IVQSNHLI
21832 52200 B 21960 I 1188					 		

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
21833	52201	В	21961	1	1414	
21834	52202	В	21962	1	1515	
21835	52203	A	21963	1652	1916	RKNPFILH*LFR*TLRQTKPDNS AGKCVKI**HTQNQRSGRSQND F*RR
21836	52204	Α	21964	1	1353	
21837	52205	A	21965		1507	MAAQDAMVTRTSSQTDCKEPG LKWAGVVLMLLLCWLMFQVL WEACQQKTGEHKTMLQMILCN KSYQQLRLVITPMYLMLGDWL SKNCGQNSTPSLGDNPPERQTQ EWSGSATFTSDGKIRLFYTDYS GKHYGKQSLTTAQVNVSKSDD TLKINGVEDHKTIFDGDGKTYQ NVQQFIDEGNYTSGDNHTLRDP HYVEDKGHKYLVFEANTGTEN GYQGEESLFNKAYYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDDLHIHAEDTL LPFYLGEKDDVTYAIKPTCWPG LDIIPSCLALHRIETELMGKFDE GKLPTDPHLMLRLAIETVAH*P AEK*HQNHSSPF*PRFFAVSLRT GSSYHCILSSH
21838	52206	A	21966	729	843	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLLINM
21839	52207	В	21967	1	1416	
21840	52208	Α	21968	1	1794	
21841	52209	В	21969	1	1171	
21842	52210	A	21970	1108	1250	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPGGASRVP K
21843	52211	Α	21971	1	2535	
21844	52212	A	21972	743	898	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPAFFQLSIE LSAT
21845	52213	В	21973	1	1605	
21846	52214	В	21974	1	1873	
21847	52215	В	21975	1	2946	
21848	52216	A	21976	1419	1498	
21849	52217	Α	21977	1	2314	
21850	52218	Α	21978	1	1215	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21851	52219	Α	21979	1	1851	PWISAPVPVDVVEGAMDSVTV
						LSFGGLMLYFCAGWPPARRWC
						FPESISCGSMERDQWWGLQVA
						KRAGLAGGQSGRTVLRERVRIE
						IASTHIALAARHSDWRCCRNGR
						YPARGPAALQNFQRYTGIQHV
						HRIGMAERMWCDRNRERHTVS
						SSGGNRLPNPGPDRSCDNLKTC
						HTSHGSVMAETAVINHKKRKN
						SPRIVQSNDLTEAAYSLSRDQK
						RMLYLFVDQIRKSDGTLQEHD
						GICEIHVA\KYVEIF\GLPFAEAS
						KDIRQALKSFAGKEVVFYRPEE
						DAGDEKGYESFPWFIKRAPSPS
						KGLYSVHINPYLIPFFIGLQN\RL
Ì						TQFRLSETKEIPNPYAM/RLYTN
						TLCQYRKPDGSGIVSLKIDWIIE
						RYQLPQSYQRVWTPGINCPEPC
						PGVWTPGINCPGAWGIHGPHGS
						AENTGPLHPGQTSAQLETSLLP
				İ		SLRSARELELAQLSPRRKQNRS
						LQNIGVTPSLRVPWPKASNVQQ
						FIDEGNYTSGDNHTLRDPHYVE
						DKGHKYLVFEANTGTENGYQG
		1				EESLFNKAYYGGGTNFFRKESQ
						KLQQSAKKRDAELANGALGIIE
						LNNDYTLKKVMKPLITSNTVTD
						EIERANVFKMNGKWYLFTDSR
21852	52220	Α	21980	3197	5376	
21853	52221	Α	21981	543	5690	RKNPFILH*LFR*TLRQTKPDNS
		ł				AGKCVKI**HTQNQRSGRSQND
						F*RRWSRSY
21854	52222	Α	21982	2241	2614	ETLTTLKTKAINTLYSKPTREQ
						KTDTKAKNLYLTKRTTAAART
				1		SSVKKAISFNRALKNAMLS*RT
						APSVS*IRHRV*NLTMRSSSSSA
]						PSPWML*A*AWLCRYCRASCGI
						SSIPTASPVTMACC
21855	52223	Α	21983	31	892	
21856	52224	Α	21984	548	855	
21857	52225	В	21985	1	1825	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21858	52226	A	21986	1	3146	MEWGKFKRLVLPSQGKSKPEQ
						RCQYRHTVGDQATLPLRSRVE
						KRNGEAEGRIYRKLWDVALLC
						CVQEVINGERKPLRALTEVKSG
						FRLREDSDKFDANDPILKDQTQ
						EWSGSATFTSDGKIRLFYTDYS
						GKHYGKQSLTTAQNVQQFIDE
						GNYTSGDNHTLRDPHYVEDKG
						HKYLVFEANTGTENGYQGEES
i						LFNKAYYGGGTNFFRKESQKL
						QQSAKKRDAELANGALGIIELN
						NDYTLKKVMKPLITSNTCDN
21859	52227	A	21987	1	3107	MNEMKREEKFREKRRERNEQI
						LQEIRDYVKRSNLHLIGVSKSD
1						GENGTELENTLQDIIQENFPNLA
						RQANIQIQEIQRMLQRYSSRRTT
						PRHIIVRFTKVEMKEKMLRAAR
						EKGRVTHKGKPIRLTADLSAET
						LQATKEWRPIFNILKEKNFQPRI
Ŀ						SYPAKLSFISEGETKYFTDKQM
						LRDFVTTRPALKELLKEALNME
						RNNRLVKSLAWASQNGFRSPA
						LLPLSLMLCAPLQEADGTNLG
						WLAHCGLLCTGDH
21860	52228	В	21988	239	2825	
21861	52229	A	21989	7533	11119	
21862	52230	Α	21990	1	3718	MEVIPEGYRLSELKKCSKWIVV
						MVARLYEYTKNQQIIHLKWVN
						HMLGCLLDRNDPLRDVDEAAV
						PVLCICSADDPVCGPPDHTLTTE
						LFHSNPYFFLLLSRHGGHCGFL
						RQEPLPAWSHEVILESFRALTEF
ŀ						FRTEERIKGLSRHRASFLGGRRR
						GGALQRREVSSSSNLEEIFNWK
						RSYTRLMAAAAGAAAAPGSRE
			1			PQDRPECGAGHPGPRYYRHPER
						WLLRPEAFLGPLRTRAPSAEDS
01050	5000	<u> </u>	2100:		608	QRERPAARSGPEM
21863	52231	Α	21991	1	697	
21864	52232	A	21992	1	978	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21865	52233	A	21993	145	1929	VSGVIEIADGSRRRKAAALTES DYRVLVGELDDEQMAALSRLG NDYRPTSAYERGQRYASRLQN EFAGNISALADAECDNLKTCHT SHGSVMAETAVINHKKRKNSP RIVQSNDLTEAAYSLSRDQKRM LYLFVDQIRKSDGTLQEHDGIC EIHVAKYAEIFGLTSAEASKDIR QALKSFAGKEVVFYRPEEDAG DEKGYESFPWFIK/RPSRGLYSV HINPYLIPFFIG\LQNRFTQFRLS ETKEITNPYAMRLYESLCQYRK PDGSGIVSLKIDWIIERYQLPQS YQRMPDFRRRFLQGFCRFNHI HQTGFSPAGANQRGPLAATLSG PGGEGQSAVARLTGEKKNHPG AQYANRLSPRVGRFINAAGTTG FPTWKAGSERNAINDDVTYAIK PTCWPGLDIIPSCLALHRIETEL MGKFDEGKLPTDPHLMLRLAIE TVAHDYDVIVIDSAPNLGIGTIN VVCAADVLIVPTPAELFDYTSA LQFFDMLRDLLKNVDLKGNSN GSQSPWMEEQIRDAWGSMVLK NVVRETDEVGKGQIRMRTVFE QAIDQRSSTGAWRNALSIWEPE
						CNEISIGVSLDQDGGSNSVLRK

2811

21866	SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
21866 52234 A 21994 1184 2229 RSSVATVALLHAVINHK/KRI SPRIVQSNDLTEAAYSLSRDQ RMILYLEVDQIRKSDFTLQV RMILYLEVDQIRKSDFTLQV GICEIHVAKYAEIFGLTSAEA DIRQALKSFAGKEVVFYRFEI AGDEKGYESFPWFIKAH AGDEKGYFIKAH AGDEKGYFFFTQGQD	NO:	1	IJOu		ŀ		deletion, \=possible nucleotide insertion)
SPRIVQSNDLTEAAYSLSRD(:	•		,			, , , , , , , , , , , , , , , , , , ,
SPRIVQSNDLTEAAYSLSRD(<u> </u>				
RMLYLPVDQIRKSDGTLQEH GICEIHVAKYAEJEIGLTSAEA DIRQALKSFAGKEVYPYRPE AGDEKGYESPPWFIKRAHSP GLYSVHINPYLIPFIGLQNRT QFRLSETKEITNPYAMRLYES CQYRKPDGSGIVSLKIDWIE QLPQSYQRMPDFRRRFLQVC NEINSKTPMILSYEKKGRR THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSQSPWM QIRDAWGSMVLKNVVRETD GKDTSLSTPAAPMVDSLIAR VMARGNAITLPVCGRDVKFT EVLRGDSVEKTSRVWSGAV GGESFIYMPLTGAAAFLSEM PERRNLKQCGYSGFAACG9 SQERKNSPRIVQSNDLTEAA* LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESPFWFIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMI YESLCQYRKPDGSGIVSLKIL IERYQLPQSYQRMPDFRRFT DHNSSPEREQNWMENEFDE EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT S2238 A 21998 188 421 21870 52238 A 21999 1 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTU PFNQRPCDNLKTCHTSHGSV AETAYNIHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHIDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SPPWFIKRSPE*RRGKFPFTQ GQD	21866	52234	Α	21994	1184	2229	RSSVATVALLHAVINHK/KRKN
GICEIHVAKYAEIFGLTSAEA DIRQALKSFAGKEVVFYRPE AGDEKGYESPPWIRRAHSP GLYSVHINPYLIPFIGLONRI QFRLSETKEITNPYAMRLYE CQYRKPDGSGIVSLKIDWIE QLPQSYQRMPDFRRRFLQVC NEINSRTPMRLSYTEKKGKG THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSQSPWM QIRDAWGSMYLKNVVRETIO GROTSLSTPAAPMVDSLIAR VMARGNAITLPVCGRDVKFI EVLRGDSVEKTSRVWSGAV GGESFIYMPLTGAAAFLSEM PERRILKQCGYSGFAACGY SQERKINSPRIVQSNDLTEAA LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESPPWFIK HSPSRGLYSVHINPYLIPFFIG NRRTGFRLSETKEITNPYAMI VYSSLCQYRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRFI DHINSSPEREQNWMENETEDI EVKGPRWYTINSSELKEHVL CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT LSR67 52235 A 21995 82 325 21868 52236 B 21996 220 11173 21870 52238 A 21998 188 421 21871 S2239 A 21999 1 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPRAGREPPFIGY PPNQRPCDNLKTCHTSHGSTV AGREVVFYRFGEDAGHEKG SFPWFIKRSPE*RRGKPPFTQ GQD							SPRIVQSNDLTEAAYSLSRDQK
DIRQALKSFAGKEVVFYRPE AGDEKGYESPPWFIKRAHS GLYSVHINPYLIPPFIGLONRI GLYSVHINPYLIPPFIGLONRI QPPGSQRIPPGSGIVSLKIDWIE QPPGSVGRMPGSGIVSLKIDWIE QPPGSVGRMPGRRRFLQV NEINSRTPMRLSYIEKKKGR(THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSQSPWM QIRDAWGSMYLKNVVRETD GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKT GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKT GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKT GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKT GROTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKT GESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAA' LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGI AFASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLVSVHINPYLIPPFIG RRFTCPRLSETKEITNPYAMI YESLCQYRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRST DHNSSPEREQNWEMERDEDE EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT S2235 A 21995 82 325							RMLYLFVDQIRKSDGTLQEHD
AGDEKGYESFPWFIKRAHSP GLYSWHINPYLIPFIGLOND GLYSWHINPYLIPFIGLOND GLYSWHINPYLIPFIGLOND GPELSTKEITNPYAMRLYE QPPGSYQRMPDFRRREICWC NEINSRTPMRLSYIEKKGKG THIFFDMLRDLLKNYDLKGF DVRILLTKYSNSNGSGSPWM QIRDAWGSMYLKNVVREITO GKDTSLSTPAAPMVDSLIAR VMARGNAITLPVCGRDVKF1 EVLRGDSVEKTSRVWSGAJ GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACG SQERKNSPRIVQSNDLTEAA* LSRDQKRMLYLFVDQIRKSD LOEHDGICEHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMI YESLQQYRKPDGSGIVSLKLIL EIRYQLPOSYQRMPOFRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTININ CKEAKNLDKRLQELLTININ NIDDLMELKNTAQELHETYT 1 1999 21870 52237 A 21997 I 1999 21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AERGENTAL SAENGRERPPGTC AERGENTAL SAENGRERPPGTC GGAPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SPPWFIKRSPE*RRGKFPFTQ GQD	ļ						GICEIHVAKYAEIFGLTSAEASK
GLYSVHINPYLIPFFIGLQNRI QFRLSETKEITNPYAMRLYE CQYRKPDGSGIVSL KIDWIE CQYRKPDGSGIVSL KIDWIE QLPQSYQRMPDFRRFILQVC NEINSRTPMRLSYIEKKKGR THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSQSPWM QIRDA WGSMVLKNVVRETD GKDTSLSTPAAPMVDSLIAR VMARGNAITLP/CGRDVKF EVLRGDSVEKTSRVWSGAVS GGESFIYMPLTGAAAFLSEM PERRNLKQCGYSGFAACG SQERKNSPRIVQSNDLTEAA* LSRDQKRMLYLPVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMF YESLCQYRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRFI DINSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEIHVL CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21869 52237 A 21997 I 1599 21869 52238 A 21998 188 421							DIRQALKSFAGKEVVFYRPEED
QFRLSETKEITNPYAMRLYES							AGDEKGYESFPWFIKRAHSPSR
CQYRKPDGSGIVSLKIDWIIE QLPQSYQRMPDFRRRIQVC NEINSRTPMRLSYIEKKKGRC THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSQSPWM QIRDAWGSMVLKNVVRETD GKDTSLSTPAAPMVDSLIAR: VMARGNAITLPVCGRDVKFT EVLRGDSVEKTSRVWSGAV: GGESFIYMPLTGAAFLSEM PERRILKKQCGYSGFAACG SQERKNSPRIVQSNDLTEAA* LSRDQKRMLYLFVQGIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLYSVHINPYLIPFIG NRFTOFRLSETKEITNPYAMF YESLCQYRKPDGSGIVSLKIL IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL* CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT S1238 A 21995 1 1599 2 1868 52237 A 21997 1 1599 2 1870 52238 A 21998 188 421 4	1						GLYSVHINPYLIPFFIGLQNRFT
QLPQSYQRMPDFRRRFLQVC NEINSRTPMRLSYIEKKGRG THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSSPWM QIRDAWGSMYLKNVVRETD GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKF EVLRGDSVEKTSRVWSGAVS GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAA' LSRDQKRMLYLFVDQIRKSD LQEHDGICEHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIKI HSPSRGLYSVHINPYLIPFIG NRFTQFRLSETKEITNPYAMI YESLCQYKRPDGSGIVSLKIL IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENFEDEI EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRINS NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRINS NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRINS S2235 A 21995 82 325 21869 52237 A 21997 1 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 1 596 MGMTDMGKSKKKSQPPSF GGAPPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AETAYINHKKRNSPRIVQSI LTEAAYSLSROQKFMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SPPW/FIKRSPE*RRGKFPFTQ GQD							QFRLSETKEITNPYAMRLYESL
QLPQSYQRMPDFRRRFLQVC NEINSRTPMRLSYIEKKGRG THIFFDMLRDLLKNVDLKGF DVRILLTKYSNSNGSSPWM QIRDAWGSMYLKNVVRETD GKDTSLSTPAAPMVDSLIAR' VMARGNAITLPVCGRDVKF EVLRGDSVEKTSRVWSGAVS GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAA' LSRDQKRMLYLFVDQIRKSD LQEHDGICEHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIKI HSPSRGLYSVHINPYLIPFIG NRFTQFRLSETKEITNPYAMI YESLCQYKRPDGSGIVSLKIL IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENFEDEI EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRINS NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRINS NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRINS S2235 A 21995 82 325 21869 52237 A 21997 1 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 1 596 MGMTDMGKSKKKSQPPSF GGAPPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AETAYINHKKRNSPRIVQSI LTEAAYSLSROQKFMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SPPW/FIKRSPE*RRGKFPFTQ GQD			ŀ				CQYRKPDGSGIVSLKIDWIIERY
NEINSRTPMRLSYIEKKKGRC							1 -
THIFFDMLRDLLKNVDLKGF							1
DVRILITKYSNSNGSQSPWM QIRDAWGSMVLENVVRETD GKDTSLSTPAMPVDSLIAR VMARGNAITLPVCGRDVKF1 EVLRGDSVEKTSRVWSGAV3 GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAA* LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESPPWFIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMM YESLCQYRKPDGSGIVSLKIL IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHIVL* CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHTYT							1
QIRDAWGSMVLKNVVRETD GKDTSLSTPAAPMVDSLIAR* VMARGNAITLPVCGRDVKFT EVLRGDSVEKTSRV WSGAVS GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAAY LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMF YESLCQYRKPDGSGIVSLKIL IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT	1						1
GKDTSLSTPAAPMVDSLIAR	ŀ						1
VMARGNAITLPVCGRDVKFT							()
EVLRGDSVEKTSRVWSGAVS GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACG SQERKNSPRIVQSNDLTEAAN LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMF YESLCQVRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENEFDEI EVGFRRWITINSSELKEHVL CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT S1868 52236 B 21996 220 1173						İ	
GGESFIYMPLTGAAAFLSEM PERRNLKKQCGYSGFAACGY SQERKNSPRIVQSNDLTEAAN LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIK HSPSRGLYSVHINPYLIPFIG NRFTQFRLSETKEITNPYAMI YESLCQYRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT							1
PERRILKKQCGYSGFAACGY							`
SQERKNSPRIVQSNDLTEAAY							
LSRDQKRMLYLFVDQIRKSD LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIKK HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMF YESLCQYRKPDGSGIVSLKIE IERYQLPQSYQRMPDFRRFI DHNSSPEREQNWMENEFDEI EVGFRRWITNSSELKEHVL' CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT							-
LQEHDGICEIHVAKYAEIFGL AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIKI HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMF YESLCQYRKPDGSGIVSLKID IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRR WVITNSSELKEHVL' CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDLMELKNTAQELHETYT NIDDLMELKNTAQELHETYT NIDDLMELKTAQELHETYT NIDDLMELKTAQELHETYT NIDDLMELKTAQELHETYT NIDDLMELKTAQELHETYT NIDDLMELKTAQELH							1
AEASKDIRQALKSFAGKEVV RPEEDAGDEKGYESFPWFIKI HSPSRGLYSVHINPYLIPFFIG NRFTQFRLSETKEITNPYAMI YESLCQYRKPDGSGIVSLKID IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21868 52236 B 21996 220 1173 21869 52237 A 21997 I 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTC PFFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							· · · · · · · · · · · · · · · · · · ·
RPEEDAGDEKGYESFPWFIKI							
HSPSRGLYSVHINPYLIPFFIGNRFTQFRLSETKEITNPYAMEYESLCQYRKPDGSGIVSLKID IERYQLPQSYQRMPDFRRFFIDHNSSPEREQNWMENEFDEIEVGFRRWVITNSSELKEHVL'CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT							
NRFTQFRLSETKEITNPYAME YESLCQYRKPDGSGIVSLKID IERYQLPQSYQRMPDFRRFFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21868 52236 B 21996 220 1173 21869 52237 A 21997 I 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTO PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							
YESLCQYRKPDGSGIVSLKID IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT							
IERYQLPQSYQRMPDFRRRFI DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISN NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21868 52236 B 21996 220 1173 21869 52237 A 21997 I 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTO PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKO SFPW/FIKRSPE*RRGKFPFTQ GQD							1
DHNSSPEREQNWMENEFDEI EVGFRRWVITNSSELKEHVL' CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21868 52236 B 21996 220 1173 21869 52237 A 21997 I 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							1
EVGFRRWVITNSSELKEHVL CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT 21867 52235 A 21995 82 325 21868 52236 B 21996 220 1173 21869 52237 A 21997 1 1599 21870 52238 A 21998 188 421 21871 52239 A 21999 1 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTC PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							
CKEAKNLDKRLQELLTRISNI NIDDLMELKNTAQELHETYT 21867 52235							•
NIDDLMELKNTAQELHETYT							1
21867 52235 A 21995 82 325							•
21868 52236 B 21996 220 1173							NIDDLMELKNTAQELHETYTSI
21869 52237 A 21997 I 1599							
21870 52238 A 21998 188 421 21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTO PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSI LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD					220		
21871 52239 A 21999 I 596 MGMTDMGKSKKKSQGPPSF GGAPPPAIGAPAGRERPPGTO PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSN LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD			-		<u>l</u>		
GGAPPPAIGAPAGRERPPGTO PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSN LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD			-				
PFNQRPCDNLKTCHTSHGSV AETAVINHKKRKNSPRIVQSN LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG' SFPW/FIKRSPE*RRGKFPFTQ GQD	21871	52239	Α	21999	1	596	MGMTDMGKSKKKSQGPPSFPP
AETAVINHKKRKNSPRIVQSY LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD			1	1			GGAPPPAIGAPAGRERPPGTGD
LTEAAYSLSRDQKRMLYLFV QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD					-		PFNQRPCDNLKTCHTSHGSVM
QIRKSDGTLQEHDGICEIHVA YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD					[AETAVINHKKRKNSPRIVQSND
YAEIFGLTSAEASKDIRQALK AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							LTEAAYSLSRDQKRMLYLFVD
AGKEVVFYRPGEDAGHEKG SFPW/FIKRSPE*RRGKFPFTQ GQD							QIRKSDGTLQEHDGICEIHVAK
SFPW/FIKRSPE*RRGKFPFTQ GQD							YAEIFGLTSAEASKDIRQALKSF
GQD							AGKEVVFYRPGEDAGHEKGYE
GQD							SFPW/FIKRSPE*RRGKFPFTQM
		1					- 1
, , , , , , , , , , , , , , , , , , , ,	21872	52240	Α	22000	1	1527	
21873 52241 A 22001 2 870	21873	52241	A	22001	2	870	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21874	52242	A	22002	1	4304	MAETAVINHKKRKNSPRIVQSN
						DLTEAAYSLSRDQKRMLYLFV
						DQIRKSDGTLQEHDGICEIHVA
ŀ						KYAEIFGLTSAEASKDIRQALKS
			E			FAGKEVVFYRPEEDAGDEKGY
						ESFPWFIKRAHSPSRGLYSNRFT
						QFRLSETKEITNPYAMRLYESL
						CQYRKPDGSGIVSLKIDWIIERY
						QLPQSYQRMPDFRRRFLQCINA
		1				GHEMTKAIAIAQFNDDSPEARK
						ITRRWRIGEAADLVGVSSQAIR
						DAEKAGRLPHPD
21875	52243	Α	22003	740	1159	SLPNLDNSAICSSSSSPTRTR*SL
						SVSAAALRRRLPSAISMTPDTL
	1]		RPNAGVC*PVSRKEGMRSSSAS
						SVSSSWSRSLPDHTREVFSTLSP
						RSTSRVNFTSRPHTGKVMALPR
						AITPTRAINESTIETAGEDNEVSS
1				1		TG
21876	52244	Α	22004	664	1844	SGNLKACHPSHGSVMAETA\VI
						NHKKRKNSPR\IEQ*KDGYQQA
						YSLSRDQKRMLYLFVDQIRKSD
						GTLQEHD\GI\CQIHGAQYR*QIS
						D*PFAEASKDIRQALKSFAGKE
				1		VVFYRPEEDAGDEKGYESFPW
						FIKRAHSPSRGLFSGHIPP\YLIPF
						FIGVKNRLTQI\RLFETKEIPHPS
				ŀ		AL\RYYESLGQDRK\PDGSSIGS
						LK\IDWIIERYQLPQSYQRMPDF
						RRRFLQHFLAHAGCPKGTQNFT
						FRMKFIIFSIKKVLPPVDTILAHD
	-			ľ		PVIHTCHLRESFGSVSPSSPKSN
				1		RLPCPEVILFHYGESWNLLRAD
						QRLIFAKSWPRASRYQQGHQD
						LFILRSDLPSQVFIRDKLMERRN
						RRTGRTEKARIWGSDGQNGQE
21077	1500.45	 	22005	3	272	PGYVEAGCPPLLLTV
21877 21878	52245 52246	A	22005 22006	1	272 528	
21878	52246	A	22007	461	831	LLNLATWRYLGTLTRGTAS\VA
210/9	32241	^	22001	1701	1001	VAGILAALRITNNKLSNHVFVF
						QGAGEAAMGIAHLLVMALRRK
					-	K\GTERTTATRKIWMVDSK\GLI
				1		CQGEGSHLN/HHEKEMFAQDH
						PEVNSLEEVVRLVKPTAM
L	<u></u>	<u> </u>	<u> </u>	<u> </u>	<u>L</u>	FEANSLEEV AND AND ALL INIT

sequence 09/540,217 codon for peptide of peptide sequence deletion, \=post 21880 52248 A 22008 3 1894 EGEDRGL APWPGRA	, /=possible nucleotide ssible nucleotide insertion)
	ssible nucleotide insertion)
21880 52248 A 22008 3 1894 EGEDRGLI APWPGRA	
APWPGRA	
APWPGRA	PRTMGAALGTGTRL
1 1 1 1 1 1 1	CGALPRWTPTAPAQ
	PARPVPLKKRGYDV
1 } 1 }	KGMAFTLEERLQLGI
1 1 1 1 1	LSQDVQLLRIMRYY
1 1 1 1	DKYIILMTLQDRNEK
	SDVEKFMPIVYTPTV
	GLTFRRPRGLFITIHD
1	ILNSWPEDNIKAVVV
	GLGDLGCYGMGIPVG
1 1 1 1 1 1	CGGVNPQQCLPVLL
	ELLRDPL\YIGLKHQ
	DDLLDEFMQAVTD
	I\QFEDFANANAFRLL
	'CMFNDDIQGTASVA
	LRITNNKLSNHVFVF
	MG\IAHLLVMALE\K
	EATRKIW\MVDF\KGL
1 1 1 1 1 1 1	NHEKEMFAQD\HPE
1 1 1 1 1	VRLVKPTAIIGVAAIA
	LRNMASF\RRAPIIFA
	AECTA\EKCYRVTEG
	GKSFLRSVDL/ESMGK
	NNA*RVPRGWQLG
1 1 1 1 1	PGHIP\DEIFLPDSRA
	EQHL\SQGRLYP\PLS
1 1 1 1 1 1	RIAIKVLDYAYKHNL
1 1 1 1 1 1	KDKEAFCKIPGSYTP
	VDSYIWAQGKAMN
21881 52249 A 22009 322 567	
21882 52250 C 22010 338 603	
21883 52251 A 22011 127 417	
21884 52252 A 22012 253 424	
	GQGDRALSRTSQWP
	.CGGSEQIPGIDIQLN
RKYHTTR	KLSTTKRFPTACWR
KKVGAFT	KIIEAMGFTGPLKYS
1 1 1 1 1 1	RALRIVY*LCGRKLD
FEGFFSK\C	CS\MPDTFNSWFLINP
	CLVRMKQEGRSGKY
MCRIIVHF	MWEDVQQRGRVM
GVNPYILK	KKNMILMTNHFYA\A
	LSDDHGLAAALWK
	CEDPRHLELLVEYVR
	S\MNGEDLLLTGEVS
	NPQSILKPHSPTYND
RGTFDGLC	GPPHGPPAGFRGTSR
1 1 1 1 1	PRTPARKWPELDLLN
1 1 1 1 1	APICVEFPLSVPQESD
LLGYRAW	'ENPLALPGCPSSLRG

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21886	52254	A	22014	282	435	GGGASPAACSGSPDSS\HSATA GTGCRRPARPPPAPPTPTPWRS RQQGSRAS
21887	52255	A	22015	3	399	IVALPFGHHHSQTSPLPNLLQTA ACLLLHFLLLSGVETGRPGPWQ PRIRARGPPAPPGPARPAKPRRR /GCRGPAREACWYGCRRPARPP PAPPTPTHGAAGSREAERASEL/ C*GARPLRLGSAGTIMVSACVP
21888	52256	A	22016	279	872	KIPIISLSLKLGQASSKIIPTPQNP HLSPGTKRDSSFQPWSLSAFIIT GQCCSYIVRWGLWKPRLILVV KKYAALSAEMLLVKVEEPSPA ACSGSPDSSSFCNCW*PAAAGR L/ASAAQRPPTPNAHKRSRQQG SRASERAS/CKGRGRYGLGALG GRGGRALGGSRWPPPLPGETLF SGCKHRRRRRGSDAAPGEEAG
21889	52257	A	22017	106	398	NAAPFSRHAAFQGFSWAVCIFR LPAERALRLFPASCLVLPGGPS GVAP\HGWPLPTTAPPLLQGVW RVEVGEHEMAEIFSSAKKEAAG ALQRPCIPVW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	-			sequence		
21890	52258	Α	22018	1	1806	MRGNTAGGPGDEKPDSTHPPW
						EGTLQEAQEMKKVVRFMWTS
		1				GGPDPQHWRGSGLYGQLVAQ
						MLHIDVAQVYMDSWWPRCST
						LAWLRFIWTAGGPDAPHWLRS
						GLGGQLVAWMLHVGVAQVYV
ŀ						DSWWPSCSASMWLSFMEHKQ
						CLTLGYEVALFLQHFLSRAFVL
		1				VWVPRKEKAQVLRQGLSLTFR
	ĺ					ENPGVCVMSSYRPWLITAAAD
						MQPSKPGGTGSYCSYWSPTDN
						TPRLAAACTWNLGSSHGSAPG
						VRTFIEAGFSPPFLAPKALALTL
						SDSDSRWGPNPVYAQPQP/PEP
						AWGPNPV*GQPQPRNRPG\PEP
						CVCSTPTAEPPWGPNPVSTPTA
						EPA WGPNPVYA/PTPTAEPPWG
						PNPV*GQPQP/REPAWGPNPVY
	,					GQPQPRNRPGS\PPQGNLYCSA
						ALGATDPSQSWHQCSQNLLPA
						DKGAKREAPW/PRPPSGEVDHQ
						VWHLHGWPLPTTAPPLLQGVW
						RVEQTPASHHRQPRPRQSSPST
						KAPPVITVNQGPASHHRQPRPR
			:			QSSPSTKAPPVITVNQGPASHH
						RQPRPRQSSPST/TGPASHHRQP
						RPRQSSPSTKAPPVITVNQGPAS
						HHRQPRPRQSSPSTKAPPVITVN
						QGPASHHRQPRPRQSSPSTKAP
						PVITVNQGPASHHRQPRPRQSSP
		ĺ				STKAPPVITVNQGPASHHRQPR
	1		ŀ			PRQSSTLTKAPPVITVNQSPASH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21891	52259	Α	22019	27	1367	RILAGPARAEAGSASDRSTGHL
		1				FPRGVGMDLCHPQPAELSSGET
		1				DELPRI*WHRKQLLEDIQKLKD
		1				EIADVFAQIDCFESAGDSRMAH
						KENELCIGRKKFNMDPRQGRW
		1				LWRARATGCGGCMGAPAW\FL
						SGEGLNKTAIGTYLGER*ERVE
						P*RGRK*RCAGVYRCVDVQAL
						R*VVLGQGPGPQGWHSPSGWG
						CRDPRYCLCNPGVFQSTGARRG
		l				VGPRAPGPLQHCQA*ISQTRPPF
						ERFVSMNRGINNGSDLPEDQLR
]		1		ļ		VREAQPTQPGPGESRRDLSLDG
1						WPFFNPDREGWLLKLGERPTD
1		1				TRRRTHTHARAGTHRCMHTTV
1		l				SRVLLGLPLPPPLPAQPTSQ/PPT
						SRC/LRLQDPE/CWSHWEIS*TE
						RGESRVVEGKHESYRISATSAE
		ŀ				ERDQWIESIR*GVPRSGDGFPSP
						SRRPCFFWVLRDSWRWHWGL
						VTLRVPSPAAPGDP
21892	52260	Α	22020	140	356	
21893	52261	Α	22021	2	1147	APARGPAVASGPSPGPAAEPVQ
						NTETAAMTTHVTLEDALSNVD
						LLEELPLPDQQPCIEPPPSSIMYQ
						ANFDTNFEDRNAFVTGIARYIE
						QATVHSSMNEMLEEGHEYAV
						MLYTWRSCSRAIPQVKCNEQP
						NRVEIYEKTVEVLEPEVTKLMK
						FMYFQRKAIERFCSEVKRLCHA
	ĺ					ERRKDFVSEAYLLTLGKFINMF
						AVLDELKNMKCSVKNDHSAY
						KRAAQFLRKMADPQSIQESQNL
						SMFLANHNRITQCLHQQLEVIP
					}	GYEELLADIVNICVDYYENKM
						YLTPSEKHMLLKVMGFGLYLM
						DGNVSNIYKLDAKKRINLSKID
						KFFK/HAAGGAPFRRHADRAGQ
						IH*DQCSL*REQVQVDVHPEQH
	<u> </u>	<u> </u>	<u></u>			QPPVQYLRADGSDPG

SEQ ID	SEQ ID NO:	Met				Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21894	52262	Α	22022	47	1581	ALVLMYLASSICMSPKRGTTCS
						CLERGSRCPTSEPPPPPEPPPGRS
						SRCRPGLGPYGGGGEKMADDP
						SAADRNVEIWKIKKLIKSLEAA
ŀ						RGNGTSMISLIIPPKDQISRVAK
						MLADEFGTASNIKSRVNRLSVL
				+		GAITSVQQRLKLYNKVPPNGLV
						VYCGTIVTEEGKEKKVNIDFEP
						FKPINTSLYLCDNKFHTEALTA
						LLSDDRQVWDFIVIGG/SVGALF
						GHTPKGNTRGGPWHKFTVGIFP
						K\KHG*EGGQ\SALRFARLRME
ŀ						KRHNYVRKVAETAVQLFISGD
						KVNVAGLVLAGSADFKTELSQ
						SDMFDQRLQSKVLKLVDISYG
						GENGFNQAIELSTEVLSNVKFIQ
						EKKLIGRYFDEISQ/DLTGKYCF
				1		GVEDTLKALEMGAVEN\LIVYE
						NL\DIMRYVLHCQGT\KEEKILY
		1				LTPEQEK\DKSPFH\DKETGQE\L
						ELIESMPLLEWFANNYKKFGA\
					i	TLEMFTDKSQKGSPV**KDLGG
		1				IGG\ILRYQSRFPGEWNTKKET
						MNFLTLMTTR
21895	52263	Α	22023	1	296	VEDERVCAGPAWERQAGQERP
						ARKVQGKAPEAEGKHGCLLDI
		1				LVCTYLAEKGLSGSCGLTEALV
		1				AFGEGSSLGRKNCRDPWGSLR
						QEPHWSDP/WSGPPERCKGKPQ
						RRRGSTDAFWTSWSART*PRRG
						SLAAVG*QRHWWPLGRAAV*E
						ERTAETPGAPSGRSLTGQTLGS
						LHV
21896	52264	Α	22024	5	309	LPDADLFCPACGAAAGRARSA
						GMRPVFAVSVARTPSPR/PPPAE
						PQFPRLPCGSIGSPFFRDIDTIQK
						FPDMLSSFFFRDIDTIQKFPDML
						VFNCCCLLNQSR
21897	52265	Α	22025	1	258	RGRSRIARSASGPRPALVRPPAR
						PPAP*TPRSWSCWSSC*PRSASA
						TGSPSA*ATDAHADSSKAMLPE
						PTSSISKFSTLQTVPFRL
21898	52266	Α	22026	119	692	PSSHSVRTATYSSLRQSAVSTRT
		1				STTTTLAFMARAGGRAGGRAR
						VGGRTPSGQCGCPACGAAAGR
		1				ARSAGMRPVFAVSVARTPSPR/
						PPPAEPQFPRRPQR\GAPAPSGG
						GASLPAPRALASSIPAQLMKPG
		1				RAKGSAGNLHPG\EGRPGARRS
						DPAGAARSLACP*IWPKQTPSP
						LASSSSFFLSETPECAH
		1	<u> </u>		<u> </u>	LAGGOST LOCATI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21899	52267	A	22027	2	246	
21900	52268	Α	22028	1	927	
21901	52269	Α	22029	251	1077	VTMGKVLAVSSLPAGPPPPPVP
[ŀ					ALVGLPPPPP*HPGFRLPPLGGG
						LGARTSVGRGLERTPGAATASA
						AGGAEDGACRCLPNPCTFEECH
						RKCKEL*MEGVKFTENKAL/SS
						KMAIQTQQSKFVKWQVDGEYR
		1				GSDFTVDVTLWNPDVLMGSGI
		l				LVAHYLKSITPCLALGSELVYH
					1	RRPGEEGTVMSVAGKYTLNNW
						LATVDVEFEASTRMQDTNVSF
						GYQLDLPKANLLFKGSVDSNW
						IVGATLEKKLPPLPLTLALGAFL
						NHRKNKLRSGFGLTIS
21902	52270	Α	22030	1	1171	GGGSGFGCAWRTGWERSPGRE
		1				QAPPPATMGNVLAASSPPAGPP
						PPPAPALVGLPPPPPSPPGFTLPP
1						LGGSLGAGTSTSRSSERTPGAA
						TASASGAAEDGACGCLPNPGTF
						EECHRKCKELFPIQMEGVKLTV
						NKGLSNHFQVNHTVALSTIGES
						NYHFGVTYVGTKQLSPTEAFPV
						LVGDMDNSGSLNAQVIHQLGP
						GLRSKMAIQTQQSKFVNWQVD
						GEYRGSDFTAAVTLGNPDVLV
						GSGILVAHYLQSITPCLALGGEL
						VYHRRPGEEGTVMSLAGKYTL
						NNWLATVTLGQAGMHATYYH
						KASDQLQVGVEFE\ASTRMQDT
						SVSFGYQLDLPKANLLFKGSVD
1					}	SNWIVGATLEKKLPPLPLTLAL
						GAFLNHRKNKFQCGFGLTIG
21903	52271	Α	22031	14	434	CYCQGILSPPPCPPFLLGSAQTS
				1	1	AICSWVRGVPAESSTSSFPKAP*
				1		VALESRGEAPTLHLPVTSRFLQ
				1		CVSSEQECVPWKP/HSSPRPRFL
						DISPSPTCVRL*APGPVPFDPHQ
						KTWRSPLGTRAQGVGPRDRDS
		١.	<u> </u>	<u> </u>		CRIVPG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>	sequence		
21904	52272	Α	22032	989	1859	VYGDSRWSLALSPRLECSGA\IS
						AH*NLCLPGSSDSPASRITGISG
						ARHHAQLI/L/YVFLVEMRFHH
						VGQAGLKLLTSGNP/HHLGLPK
		İ				CWDYSRKPPPRPPRPII\FSFFFL
		İ				RWSLALVAQAGMQWDDLSSL
						QPLPPGFK*FSCLSLLRSWDYG
ļ						CPPPRLANFVCVCVCVLV/VDG
		ļ			}	GFTMLARLGLELLTSGDLPPFG
						LPKCLGFTGMSHCARPIFFF\FE
						MESCFATQAGVQWCDLSSLQP
						LPPRFK*FSCLSLLSSWDYRHM
						PPCLANF/CFGIFSRDRVSSCWP
						GRSQTPDLK
21905	52273	Α	22033	157	376	SHQDFMRTAANLLQTGWCTGS
	1					NTGAGARWGLCRASLSRCRCA
						APSAPCSGSRCS*RSWKGTHPS
						LPPRCRGRS
21906	52274	Α	22034	298	399	
21907	52275	Α	22035	1	185	
21908	52276	Α	22036	311	516	WKAERRTRHLLQTGWCTGSNT
						GAGARWGLCRASLSRCRCAAP
						SAPRSGSRCS*RLWKGTHPSLPP
		<u> </u>				RCR
21909	52277	Α	22037	1	1487	MMGHSSAIPLTATPGELKGQSP
						TKMPDPELGCQGAKSQGCSRN
İ						ARHQKARSMPLQDQHLALAIL
						LELAVQRGTLSQMLSAILLLLQ
						LWDSRAQETDNERSAQGTSTL
						LLSLLQTFQSIICSKDTPPSEGN
						MHLLSGPLSPSESFLRESFFTVQ
						NCRNNEEVTLICKADLENHNK
,						DGGFWIVIDEKVYDIKDFQTQS
ļ						LTGNSILAQFAGENPVVALEAA
						FEFEVTRESMHAFCVGQYLEVR
						LYALSDAEDGRGHPLMASVIHL
						LWRLQTSQIHYSYNEEKDEDH
						CSSQWAHLPANRDCSHRWALG
						DHSQAFLQAIADNNIQDHNVKT
						HQEQGRSYKEVCTPVIERLRFL
						SNELRPA VGNDLSIISEFKLLSSL
						PRWRRIAQKIIRERRKKRIPKKP
						ESTADEEKIGNEESDLEEACILP
						HSPINVDKRPIAIKSPKTITSENP
						LG/REPGEHPASPLPPDDAQHA
						HPAAQRKQPRPPAQFRHAGPHS
			<u> </u>		<u></u>	DGTAPDWPQL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21910	52278	Α	22038	263	375	AGPRRMRRISAWAVSTPLTDSP
						SPGHQPA/GEVAGAGSF**HPP/P
						PCPSRYCPQMTPPPPPPAPPPAL
						PPPPPPAPASPPLPPAPSPPPHAP
						PQALPPPPALPPPPPASPPLLPPA
						LSPPLPAPP/PPSAPPASPPPAPPP
						PPAPPPSPPPAPPPSAPSSPAPLPP
				-		APASPP\SPAPPPPAPPPAPHPPSP
		ļ				PAPPCPPLPRPRGPSVPQEKVHQ
		İ				V*IPVP\PAGEGGPAPGRIWPQG
		ŀ		į		WRRAGKPPTWEEALAAFPPWT
l						ALPPASAVPAQPPRAH*VAPAL
						*WALHRRSDSDSDSSNLSPRIPG
						TGGQRRQPR*DS*APACSEVAG
		ŀ				AGSF
21911	52279	Α	22039	1	2001	11001
21912	52280	A	22040	1846	7725	PEPFCHRAAHWSNQYCPFQFQ
21712	32200	l	22010	10.0	1,123	DTDSPRQTLTVLIKDNGEPSLST
1						TATLTVSVTEDSPEARAEFPSGS
						APREQKKNLTFYLLLSLILASV
						GFVVTAFGVIIFKVYKWKQSRD
						LYRAPVSSLYRTPGPSLHADAV
						RGGLMSPHLYHQVYLTTDSRR
						SDPLLKKPGAASPLASRQNTLR
	1					SCDPVFYRQVLGAESAPPGQIR
						YPVPEESQEGTFVGNVAQDFLL
						DTDSLSARRLQVAGEVNQRHF
						RVDLDSGALLIKNP
21913	52281	Α	22041	108	340	PQPSQVMSFHGDSGSCRRWAT
21713	32201	``	22041	100	3.0	AGT*ATSPPCSSSIFLRRMHSGH
						WCSCWPVRGTPCRDFTAQMAG
						PSRGSKTNRSMW
21914	52282	A	22042	275	752	EKTHKTCRCGRGCGKRGTL*H
121717	32202	l' `	22072	2,3	1,32	CWWECNLLQPLWKMVW\KWL
						KELKVELLCDLAVPL/LGINPEE
						KKSLYEKDTCT/R/LFMAAQLAI
						AKKWNQPKCPSINKWLQKLW
						YIYMMEYYSAIKRN\LTAFAAT
						WMRLETIILSQGDPLESTCSRSQ
						ANRQEITSVPKA
21915	52282	A	22043	25	202	DAEAWAMELADVGA*ASSQG
21913	52283	A	22043	23	202	APQPEEPAASLGVNEKGGAGG
						RSGRRPLGRDRPRPTRLL
		L	<u>I</u>			ROURTLURDRYRFIRLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21916	52284		22044	<u> </u>	2290	MEKLGVEPEEEGGGDDEEDAE
21910	3220.	` `		•	1230	AWAMELADVGAAASSQGAPQ
						PEEPAASLGVNEKGVHDQVLPT
						PNASSRVIVHVDLDCFYAQVE
						MISNPELKDKPLGVQQKYLVV
						TCNYEARKLGVKKLMNVRDA
						KEKCPQLVLVNGEDLTRYREM
						SYKVTELLEEFSPVVERLGFDE
						NFVDLTEMVEKRLQQLQSDEL
						SAVTVSGHVYNNQSINLLDVLH
						IRLLVGSQIAAEMREAMYNQL
						GLTGCAGVASNKLLAKLVSGV
						FKPNQQTVLLPESCQHLIHSLN
						HIKEIPGIGYKTAKCLEALGINS
						VRDLQTFSPKILEKELGISVAQR
						IQKLSFGEDNSPVILSGPPQSFSE
						EDSFKKCSSEVEAKNKIEELLAS
						LLNRVCQDGRKPHTVRLIIRRY
						SSEKHYGRESRQCHIPSHVIQKL
						GTGNYDVMPPMVDILMKLFRN
						MVNVKMPFRLTLLSMCFCSLK
						AVNTATTRLIDYYLMPTISITSR
						SGKHSFKMKDTHMEDFPKDKE
						TNRDFLPSGRIESTRTRESPLDT
						TNFSKEKDINEFPLCSLPEGVDQ
						EVFKQLPVDIQEEILSGKSREKF
						QGKGSVSCPLHASRGVLSFFSK
						KQMQDIPINPRDHLSSSKQVSS
						VSPCEPGTSGFNSSSSSYMSSQK
						DYSYYLDNRLKDERISQGPKEP
						QGFHFT\NSNPA\VSAFHS\FPNL
						QSEQLFSRNHTTDSHKQTV\AT
21917	52285	Α	22045	2	141	
21918	52286	С	22046	278	489	
21919	52287	Α	22047	130	426	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21020	52200	L	22048	357	1579	QRRPRPFPSQGISMTECFLPPTSS
21920	52288	Α	22048	337	1379	f = '
						PSEHRRVEHGSGLTRTPSSEEIS
1						PTKFPGLYRTGEPSPPHDILHEP
						S*SNGLMDEKDHGKKKGNFKK
						KEKRTEGYAAFQEDSSGDEAE
					,	KLMPRFEEAYGRTTETEKVQEF
İ						QRLLKELPEGNYLLISLFVVHM
			i:			DHVITKELEAKMNIQNIYIVVSP
					·	TVQISNLVLYVFFTHVQELFGN
						VVLKQVTKTRRWSNVATMPTL
						PETQAGFKEEIRIQELLLNCLHQ
1						DLQGGIKDLSEEERLWEEKEIL
ŀ						TKQEELLGMEQFLRRQIASEKE
						EIECLRAETAEIQSHQQQGLSET
						EEYSSESESEDEEELQIILEDLQR
						QNEELEIKNNHLNQAIHEESEA
						VIEPRMQPWLLQLQPDRAKQQ
						AQEDEEPEWRGGAMQTPKNGI
21921	52289	Α	22049	182	2212	VTEKTSSLPSQGISMTECFLPPT
	ŀ					SSPSEHRR\VEHG/SAGLTPGPPA
						SEEIQPLPKFPGLY\RTGEPPHPP
						P*HPSHEPSLM*CPDDEKDHGK
						KKGKFKKKEKRTEGYAAFQED
						SSGDEAESPSKMKR\SKGIHVFK
						KPSFSKK\REKDFKIKEKPKEEK
						HKEEKHKEEKHKEKKSKDLT\A
						ADVVKQWKEKKKKKKPIQEPE
						VPQIDVPNLKPIFGIPLADAVER
						TMMYDGIRLPAVFRECIDYVEK
						YGMKCEGIYRVSGIKSKVDELK
						AAYDREESTNLEDYEPNTVASL
						LKQYLRDLPENLLTKELMPRFE
						EACGRTTETEKVQEFQRLLKEL
1						PECNYLLISWLIVHMDHVIAKE
						LETKMNIQNISIVLSPTVQISNR
						VLYVFFTHVQELFGNVVLKQV
						MKPLRWSNMATMPTLPETQAG
						IKEEIRRQEFLLNCLHRDLQGGI
						KDLSKEERLWEVQRILTALKRK
						LREAKRQECETKIAQEIASLSKE
						DVSKEEMNENEEVINILLAQEN
						EILTEQEELLAMEQFLRRQIASE
						KEEIERLRAEIAEIQSRQQHGRS
						ETEEYSSESESESEDEEELQIILE
						DLQRQNEELEIKNNHLNQAIHE
						EREAIIELRVQLRLLQMQRAKA
						EQQAQEDEEPEWRGGAVQPPR
						DGVLEPKAAKEQPKAGKEPAK
21922	52290	В	22050	34	322	
	1			l	I	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
21022	52201	<u> </u>	22051	127	1546	QMLGGRPPPRWLQLAKARALA
21923	52291	Α	22051	137	1340	()
						RRRPPPSLHLTLPTALSLKS/RTP
		l	ļ			VPKSIPISETPNIPPVSVQPPASIG
		l				PPLGVPP/PEPSHGDDQPRPG\PL
						PIFMEQQIMQQIRPPFIRGPP\PM
•		l				PPTPTAPCPTPCFPASGPRPVAP
		l				ETWAPLPAPCTGPCY\PPHPPPE/
		l	<u>.</u>			PPPP/SPGNPPGLLPPPPPGAPLPS
						LPFPPVSMMPNGPMPVPQMMN
						FGLPSLAPLVPPPTLLVPYPVIV
				•		PLPVPIPIPIPIP\PSATPSPPTGSP
						ATGRTSFRTPLATPRRRAASQA
						DTPCPPGTPSRARPSPRTRPPAA
						RARP*AWRPRPPSMAGARWWT
		l				*RGAPA\PPGPPGRGRPARLPRR
		İ	:			AAGPAGRRHRPDRGPPSPAAQR
						DPPRAA/PRTSRRSASRAPRSAG
						PAAAAGTATAARPPPATQARA
						PRRAPRRPRPATSS*TARAAPPP/
						SGAKSAEPPPEQPPPPPPPAPPK
						KLLSPEEPAVMRARVGQGE
		ŀ				
21924	52292	A	22052	762	2049	RSSFSMGADLLACST/SGLGNM
						PY\DLLRGMEPTLGCSFEDLAL
		İ				A*PGGITQE\KINEMRVAPEQQ
						MI\ADIHFMIAAGQDLDWIDAQ
						GATLLHIAGANGYLRAAELLLD
						HGVRVDVKDWDGWEPLHAAA
		ŀ				FWGQMQMAELLVSHGAN\LNA
						RTSMDEMPIDLCEEEEFKVLLL
						ELK\HKHDVIMKSQLRHKSSLS
						RRTSHRQAS/SVGKVVRRTQPV
						GTGPNL\YRKEYE/GEEAILWQR
						SA\AEDQRTSTYNGDIRET\RTD
						QENKDPNPRLEK\PVLLSEFPTK
						IPRGELDMPVENGLRAPVSAYQ
						YALANGDVWK\VHEVPDYSMA
	:					YGNPGVADATPPWSSYKEQSP
						QTLLELKRQRAAAKLLSHPFLS
						THLGSSMARTGESSSEGKAPLI
						GGRTSPYSSNGTSVYYTVTSGD
						PPLLKFKAPIEEMEEKVHGCCRI
21925	52293	Α	22053	2	306	SKEAFAIGLGGGSASGKTTVAR
41743	JLL73	^	22033	-	500	MIIEALDVPWVVLLSMDSFYK/
						NAFDFDLIISTLKKLKQGKSVK
						APIYDFTTLSRKMDWKTLYGA
21026	52204	Α.	22054	2	270	NVIIFEGIMAFADKTL
21926	52294	Α	22054	2	379	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>		·	<u> </u>	
21927	52295	Α	22055	352	1227	GDVSPQLSPRQKSGPLCTSCCA
		1				WPVCSPHSDAQPSCGEGGLWE
		l				KTPFLSVPHRFVPRAALASAHQ
			,			CHPLPRTLSVLKSTPQVRGMHT
						IIRDKETSRDEFIFYSKRLMRLLI
						EHALSFLPFQDCVVQTPQGQD\
						*RITGVSILRAGETMEPALRAVC
			_			KDVRIGTILIQTNQLTGEPELHY
		ŀ				LRLPKDISDDHVILMDCTVSTG
						AAAMMAVRVLLDHDVPEDKIF
		ŀ				LL\SLLMAEMGVH\SVAYAFPR
		ŀ				VKKSSPRRWDKAGQLTFFRIIP
		l				GIGNFGDRYFGTDAVPDGSDEE
						EVAYTG
21928	52296	A	22056	72	2746	FLVETGLHHVGQAGLKLLTSN
		-		-		DPPASASQSTGITGMSHHAQPR
		1			}	QSSYVPPMSLPASAMPVEGVG
					=	GDALWAGHASGYLGGGQLWA
		l				TSEYIPLCSSNAESLDRLLPPVG
		İ				TGRSPRKRTTRQCKSEPPLLRTS
		ŀ				KRTIYTAGRPPWYNEHGTQSKE
		İ				1 - 1
						AFAIVPVEQEKEAGQPQAAAA GWAPLGPICAGSLPVPHRGSPL
						PPAGLGGGSASGKTTVARMIE
						ALDVPWGGSSLSMDSFYKVLT
						EQQQEQAAHNNCNFDHPDAFD
						FDLIISTLKKLKHGKSVKVPIYD
		1				FTTHSRKKDWKTLYGANVIIFE
				}		GIMAFADKTLLEGPGPAPCTGP
						EDSIIRHIAAPSLHKASSRANYC
						VLSMGTDPKLATSPIPQCVGAS
						YPKQRCPGEVALARPTDQAAS
						VQGLVPVSPCPLLASGHSLARC
						LHQLLDMKIFVDTDSDIRLVRR
					•	LRRDISERGRDIEGVIKQYNKFV
						KPSFDQYIQPTMRLADIVVPRG
						SGNTVGHRP*LLQHVHSQLEEV
						SGVPRPHGLAPSPSTHFSAATTP
						SWPVHLIPLRSLSPASS\SSQPAQ
						LCQQELGNCSKKLGLGRGRIPL
		1				SLPKGRHAGGAALASAHQCHP
						LPRTLSVLKST\PQVRGMHTIIR
	1					DKETSRDEFIFYSKRLMRLLIEH
						ALSFLPFQDCVVQTPQGQDYA
						GKCYAGKQITGVSILRAGETME
						PALRAVCKDVRIGTILIQTNQLT
21929	52297	A	22057	61	341	
21930	52298	A	22058	1	1086	
21931	52299	A	22059	1	498	
21932	52300	A	22060	104	3395	
21933	52301	A	22061	1	1653	
L	122201	ī.,	10.	1	1 - 200	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21934	52302	A	22062		1717	
21935	52303	Α	22063	172	13375	EDLFKCIMAHAASQLKKNRDL
						EINAEEEPEKKRKHRKRSRDRK
			ļ			KKSDANASYLRAARAGHLEKA
						LDYIKNGVDINICNQNGLNALH
						LASKEGHVEVVSELLQREANV
			į			DAATKKGNTALHIASLAGQAE
		ŀ				VVKVLVTNGANVNAQSQNGFT
		l				PLYMAAQENHLEVVKFLLDNG
						ASQSLATEDGFTPLAVALQQGH
						DQVVSLLLENDTKGKVRLPAL
						HIAARKDDTKAAALLLQNDNN
						ADVESKSGFTPLHIAAHYGNI
21936	52304	Α	22064	1	360	
21937	52305	A	22065	1	2577	
21938	52306	В	22066	323	3154	
21939	52307	Α	22067	93	791	IRMSYAEKPDEITKDEWMEKL
						NNLHVQRADMNRLIMNYLVTE
						GFKEAAEKFRMESGIEPSVDLE
1						TLDERIKIREIDT*KGQIQEAIRL
		1				*FNSLHPEL/LWDTNRYLYFHF
		1				ASNSILIPS*FRQ\GKTRGRRWG
	İ					VAQTQLAD\RGEESRECLTEME
						RTLALLAFDSPEESPFGDLLHT
						MQRQKVWSEVNQAVLDYENR
						ESTPKL\AKLLKLLLWAQNELD
		<u> </u>				QKKVKYPKMTDLSKGGD
21940	52308	Α	22068	171	426	
21941	52309	Α	22069	5	655	LLQPSVQLPNGTANTSLSFAAV
						SAVSPPPCRTSTATTLPPPMPSF
ļ				ł		FCVFPSPSMSPSPSEFLSCIASVS
						RVHSLSSSSSGSSSTASSLNFSAI
						MGSSSATASWVLSTTSTPPCPS
		1				ALPSSPAQES*SLAASSSAWLSE
]		HPSFSSFTLSWASGAFCDEVSLS
						FSGSKLSFLSSKGDCGIPFI*TSA
		1		1		CPSSLDFSPCFSSSTSKSSSYSEI
L						GKSPQPI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21942	52310	Α	22070	87	1180	NQASLEVLPSPVDSPDSIPSLPP/
,				}		SDNPNRNFSAAPVSCLPSPLPVA
						ASWPSDPSSCSCPSPEVFRSRAP
						GVPTPASVVYVTVISSAAGTSSP
				1		LRANSSTTTFPAAAVFPTNSSC
						WCQVSSSATSEKEEALFAIVFSS
						AVSEPANTSLSSAAVSAVSPPPC
				<u> </u>		RTSTATTLPPPMPSFFCVFPSPS
				<u> </u>		MSPSPSEFLSCIASVSRVHSLSSS
						SSGSSSTASSLNFSAIMGSSSAT
						ASWVLSTTSTPPCPSALPSSPAQ
						ES*SLAASSSAWPVAGISPSGAC
						TFPAGSASGAAKAPSPSWRCPS
l						FRALFSLLDSSSLSLCSVSLASL
	1	:				SLSPPPSFLRAVMFPATFKSRRC
	:					FKYGNHCLFSDLLWLEVCAL
21042	52211	_	22071	841	1052	 IGEVLLDNVLKSVF*LGSVLPIT
21943	52311	Α	22071	841	1052	SLKLLP/SGSKRDPAKMSHSHP
:						AG*VTTGAFPRCGRWRLQPAA
						PPPPS
21944	52312	A	22072	817	866	HANPVSPPTSWPKE*SWP*CSG
						**RTFQSVSFAGGRRRGSWLKP
						PPATAWKSSSCHLPSGVRMAHF
						CRIPFGPRGRDFADFGTTIKQDF
						RLLGQTSVDRLLQLSQGQAVK
						GNQLLPVSLVKRKTTLAPNTQT
						ASPRALADSLMQLARQVSRLES
						GQ
21945	52313	Α	22073	2	3238	TTFPAAAVFPTNSSCWCQVSSS
	-					ATSEKEEALFAIVFSSAVSEPAN
						TSLSSAAVSAVSPPPCRTSTATT
	ļ					LPPPMPSFFCVFPSPSMSPSPSEF
						LSCIASVSRVHSLSSSSSGSSSTA
						SSLNFSAIMGSSSATASWVLSTT
						STPPCPSALPSSPAQES*SLAASS
1						SAWPVAGISPSGACTFPAGSAS
						GAAKAPSPSWRCPSFRALFSLL
		l				DSSSLSLCSVSLASLSLSPPPSFL
						RAVMFSSDFSNPGVASNMGTT
21046	50014		22074	0.0	1071	VSF
21946 21947	52314 52315	C	22074	73	4374 616	VKESLTQKSKRFQNKLINRRG
41741	32313	^	22013	1'3	010	WERPRSGASWTQRRALGGRGC
						VDRWLGII/WPHRIPGARLRGFP
				1		ALNPTPQTAPSPSPDPRPRPP/CP
						SDPGPRTEAQCSL\PPPP\PSPRH
						LDPGPEPPTPLTQQGQKGTRAP
			1			RPRCPRSQK*ATSGPAVRRREA
						RGSASLRTRAPSRDSTHAGIQG
						NVALQP
Щ	<u> </u>	1	<u> </u>	<u>.</u>	I	I

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21948	52316	Α	22076	1	421	GTVVMGLPKYFISLSTKFPLSFS
						LSADELHIASLFLWPRSKPFGVI
						QRSCASQLNALPEVLKNPGDPD
				}		KMLRFAESPRNLPAGVL/GDIW
	1					TDGKGSVSLQEQVYHFPLIRPT
		Ì				PGS*ARASFLSCG*EEFRRHGLT
						FSSTRE
21949	52317	В	22077	211	2041	
21950	52318	Α	22078	200	1525	IMASAHLATSAFCTRSSSSGRSG
						SCKRGSWDSEQALIRAWTTLPS
						SLHAADGSFAGGRGIPSKRWS
						WRAGQQSSLCLIMQTSSNCGNS
						FQLVSEGASWRGLPHCSCAELQ
						DSLNFSYHPSGLSLSVRPSSPGN
!						SPKEQPFSQVLRPEPPDPEKLPV
						PPAPPSKRHCRSLSVPVDLSRW
	ļ					QPVWRPAPSKLWTPIKHRGSGG
	ļ					GGGPQVPHQSPPKRVSSLRFLT
						SSQCLFSMCPSSQTLQPSFLQPG
						PGP\SSSRPCAASPQSGSWESDA
		1				ESLSPCPPQRRFSLSPSLGPQAS
İ						RFLPSARSSPASSPELPLAT*G/L
						SATFPESRSQPCDLDARKTGVK
						RRHEEDPRRLRPSLDFDKMNQ
						KPYSGGLCLQETAREGSSISPP
						WFMACSPPPLSASCSPTGGSSQ
						VLSESEKEEEGAVRWGRQALS
1						KRTLCQRDFGDLDLNLIEEN
21951	52319	Α	22079	1	117	
21952	52320	Α	22080	3	632	
21953	52321	Α	22081	1	448	MQSETETVHLFIPALSVGAIIGK
						QGQHIKQLSRFAGASIKIAPAEA
						PDAKVRMVIITGPPEAQFKAQG
						RIYGKIKEENFVSPKEEVKLEA
						HIRVPSFAAGRVIGKGGKTAST\
						*QNLCSGMKGKALKAKPSTYN
						CVGCGWSPRPRCSLN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	IL.	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21954	52322	A	22082	233	2096	RREDEGNKTSWIHPSPWVVLIF RFFSLFFFKQPLFTMNKLYIGNL SENAAPS\DL\KSIFKDAKIPVSG PFLVKTGYAFVDCP\DESWA\LR AIKAFSGKIELHGKPIEVEHSVP KRQRIRKLQIRNIPPHLQWEVL DSLLVQYGVVESCEQVNTDSET AVVNVTYSSKDQARQALDK/L LDKLNGFQLENFTLKVAYIPDE MATQQNPFLQPRGRRGLGQRG S\SRQGSPGSVSKQKPCDSPLRL LVPTQFVGAIIGKEGATIRNITK QTQSKIDVHRKENAGAAEKSIT IPSTPEGTSAACKSILEIMHKEA QDIKFTEEIPLKILAHNNFVGRLI GKEGRNLKKIEQDTDTKITISPL QELTLYNPERTITVKGNVETCA KAEEEIMKKIRESYENDIASMN LQAHLIPGLNLNALGLFPPTSG MPPPTSGPPSAMTPPSPQSEQSE SETVHLFIPALSVRALISKQGQH IKQLSRFAGASSKIAPVEAPDAK VRMVMIAGSPEARFKAQGRIY GKIKEENFVSPKEEVKLEAHIR VPSFAAGRVIGKGGKTV\NELQ
						NLSSAEVVVPRDQTPDENDQV VVKITGHFYACQ\VAQRKI*EFL TQVRQHQQQKALQSGPPQSRR

SEQ ID NO:	SEQ ID NO: of peptide sequence	4	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21955	52323	Α	22083	8	1675	TASCRLAMALHVPKAPGFAQM
1						LKEGAKHFSGLEEAVYRNIQAC
1						KELAQTTRTAYGPKGMNKMVI
1						NHLEKLFVTNDAATILRELEVQ
1						HPAAKMIVMASHMQEQEVGD
						GTNFVLVFAGALLELAEELLRI
1						GLSVSEVIEGYEIACRKAHEILP
ĺ						NLVCCSAKNLRDIDEVSSLLRT
1				-	ļ	SIMSKQYGNEVFLAKLIAQAC\
İ			:			YLFFLIPAISMLITSEFVK/FLGS
ĺ						GISSSSVLHGMVFKKETEGDVT
ĺ						SVKDAKIAVYSCPFDGMITETK
Í						GTVLIKTAEELMNFSKGEENLM
						DAQVKAIADTGANVVVTGGKV
						ADMALHYANKYNIMLVRLNSK
						WDLRRLCKTVGATALPRLTPP
						VLEEMGHCDSVYLSEVGDTQV
			ļ			VVFKHEKEDGAISTIVLRGSTD
			· -			NLMDDIERAVDDGVNTFKVLT
					İ	RDKRLVPGGGATEIELAKQITS
İ						YGETCPGLEQYAIKKFAEAFEA
			·			IPRALAENSGVKANEVISKLYA
1	1					VHQEGNKNVGLDIEAEVPAVK
						DMLEAGILDTYLGKYWAIKLA
ĺ						TNAAVTVLRVDQIIMAKPAGGP
1						KPPSGK*NWLNFYCR
21956	52324	Α	22084	175	278	

2830

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	,
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21957	52325	Α	22085	208	1976	GPSPLKEHFSGLEETAYM\NILT
						CMELAQTTRTAYGPNGMNKM
						VINHLEKLLVTNDAATILRELE
					Ì	VQHPAAKMIVMASHMQEQEV\
						GDGTNF\VLVFAGSFPVNYLKN
						L\WRIGLSVSEVIEGYEIACRKA
						HEILPNLVC\CSAKNLRDIDEVS
						SLLRTS\IMSKQYGLMKVF\LAK
						PYLLQACRIYFSWIPAISMLITSE
						FVK/FLGSGIRSS*VLAGMVFK
						KET\EGDVTSVKDAK\IAVYS\C
}						PFDGHG*QET*GEQCLDKRTAL
						KDLDGIFR*GEQENLHG/AATSP
						KAIADTGANVVVTGGKVADM
						ALHYANKYNIMLVRLNSKM/W
						DLRRLCKT\VGATALPR\LTPSL
						SLKEMGHCD\SVYLSEVGDTQ\
						VVVFKHEKEDGAISTIVL\RGST
						DN/L*LDDIER\AVDDG\VNTFQ
						GSLQRDKRLVP\GGGSNRNLN*
						PKQITS\YGET\CPGLEQYA/ILK
						KFA*GHLKAIPPAHWQKYSGE
	,					NSGVKAQWK*SPKLYAVHQEG
						\NKNV\GLDIEAEVP/SLVKDML
						EAGIPRYLTLGKYWAIPLATNA
						AVTVLRVDQHTLV*FFFMQIIM
						AKPAG/GGPKPPKWGRKTWGD
						DQNGLKIGFNFYCR
21958	52326	Α	22086	1	346	PF*CTPSCLWAF*TARSAQTWS
21730	32320	`	22000	•		APSTRSWCSEWPKSLPSPTWLR
						*FRKVSSYSWSQSWSSPTASTD
						SSRPPHSLGASCHPGHQRSLMP
						PSR*RATSTAYASLSSSHRPNTQ
	1					PSPS
21959	52327	A	22087	3	484	RSSQPRVGLCGKSLQP/SSEGSC
21739	32321	l' `	22007	ľ	104	HLQGWCSFPRSVSSIMAEA*RT
]						*PRVSSSMHGRGPTQNREATKK
	}					ASSSETSV/QVTAQEAQAEPGA
						AGVAG/SPASAPQDSAGACWPA
						-
1						QVL*GGSDRSCRGSHPGNTEPP
						SVLRTGTAPGA*G/SSPFSPTTQS
21000	52220	D	22000	101	250	GSCCFPFSS
21960	52328	В	22088	181	359	

APWRSSNGQVPTLT*PGSTLPP G'SGKSPPI/NASGGPARAKPHEAN VHVSPPASGGSPEDV4ELPAAV AVRA*QAGSQRLPAPVCVGE MGTLCPGEPADVHQVVCGLSE PRAPHRPGGPQARAHGVPSGQS LCPAQPG*DDS/AKVSSYSWSQ SWSSPTASTDSSRPPIISLGASCC PGHQRSLMPPSR*RATSTAWRA RTASTPRCLGPRPAPWSCASLS SHRPNTQPSSPSTSVRRARLATT SSHGWALAPWTPMAPTQPTTW TRWGKTVSGRQMNTWRRPWS TCARYSGSAKRSSGSSHSPWAP PRRMEKSNSPTASWVRTDSS LRPWGGTRSSMGCGGLKLSTR GTRSCSPSGAMRSPAWSAHSLC CRLPSTTDLQDRWRLCVPGMT SSAASVATTSONLGWPAGTC*A LWGGGRWPATPAAIQAQPALP GQLPTR WSRCCWPSSWPLGWPASWPASWPASWPLGWPASWPLGWPASWPLGWPASWPLGWPASWPLGWPASWPLGWPASWPLGWPASWPASWPASWPASWPASWPASWPASWPASWPASWPAS	SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21962 52330	21961	52329	A	22089	290	2014	G/SGKSPP/VASGGPARAKPHRA VHVSPAPSGGSPGDVAELPAAV AVRA*QAGSGQRLPAPVCVGE MGTLCPGEPADVHQVVCGLSE PRAPHRPGQPQARAHGVPSGQS LCPAQPG*DDS/AKVSSYSWSQ SWSSPTASTDSSRPPHSLGASCH PGHQRSLMPPSR*RATSTAWRA RTASTPRCLGPRPAPWSCASLSS SHRPNTQPSPSPTSVRRARLATP SSHGWALAPWTPMAPTQPTTW TRWGKTVSGRQMNTWRRPWS TCARYSGSAKRSSGSSHSPWAP PRMRMEKSNSPTASWVRTDSS LRPWGGTRSSMGCEGLKLSTR GTRSCSPSGAMRSPAWSAHSLG CRLPSTTDLQDRWRLCVPGMT SSAASVATTSQNLGWPAGTC*A LWGGGRWPATPAA\QAQPALP GQLPTRWSRCCWPSSWPLCSAS GPSHARCCSPWAMSSTPLP*HC *PSGGSCTSPEGVSCLQSRLEGF ATQPHPWAERTWEAPPGGNTV ILGLLERGSCSRRGIWRKRNQE
21964 52332	21962	52330	A	22090	363	487	
21965 52333 A 22093 1 206 LFPAFSDSRECKLMKKLLEA\H EEQNVD\SYTEA\VKEYDSIFPG WTQWLTTL\LLRIKKTIQG\DEK DLR 21966 52334 A 22094 128 1062 KPFEAAMYNSGKEAEAMALLA EAER\KV\KNSQSFFSGLFGGSS KIEEACE\IYARAANMFKMAKN WSAAGNAFCQAAQLHLQLQSK HDAATCFV\DAGNAF\KKADPQ GGPFNCFDCVAFEIY\TDMGR\F TIAAKHHISIC*DLLRQKLVDIE KAIA\HYRALADYYQGEESNS\S ANKCLLKVAGYA\ALLEQYQK\ AIDIYEQGWGPNAHWTSPPPSK Y\SAKDYF\KAALCHFCMDML QA\KLAF\QKYEELFPVFSDSRE CKLMKK\LLKA\HGEQNVD\SY TE\SVKGITDSISRL\DQWL\TTH CLRI\KKTIRGD	21963	52331	A	22091	234	282	
EEQNVD\SYTEA\VKEYDSIFPG WTQWLTTL\LLRIKKTIQG\DEK DLR 21966 52334 A 22094 128 1062 KPFEAAMYNSGKEAEAMALLA EAER\KV\KNSQSFFSGLFGGSS KIEEACE\IYARAANMFKMAKN WSAAGNAFCQAAQLHLQLQSK HDAATCFV\DAGNAF\KKADPQ GGPFNCFDCVAFEIY\TDMGR\F TIAAKHHISIC*DLLRQKLVDIE KAIA\HYRALADYYQGEESNS\S ANKCLLKVAGYA\ALLEQYQK\ AIDIYEQGWGPNAHWTSPPPSK Y\SAKDYFF\KAALCHFCMDML QA\KLAF\QKYEELFPVFSDSRE CKLMKK\LLKA\HGEQNVD\SY TE\SVKGITDSISRL\DQWL\TTH CLRI\KKTIRGD	21964	52332	A	22092	742	990	
EAER\KV\KNSQSFFSGLFGGSS KIEEACE\IYARAANMFKMAKN WSAAGNAFCQAAQLHLQLQSK HDAATCFV\DAGNAF\KKADPQ GGPFNCFDCVAFEIY\TDMGR\F TIAAKHHISIC*DLLRQKLVDIE KAIA\HYRALADYYQGEESNS\S ANKCLLKVAGYA\ALLEQYQK\ AIDIYEQGWGPNAHWTSPPPSK Y\SAKDYFF\KAALCHFCMDML QA\KLAF\QKYEELFPVFSDSRE CKLMKK\LLKA\HGEQNVD\SY TE\SVKGITDSISRL\DQWL\TTH CLRI\KKTIRGD	21965						EEQNVD\SYTEA\VKEYDSIFPG WTQWLTTL\LLRIKKTIQG\DEK DLR
1010/E 150305 IA 100005 II 11000							EAER\KV\KNSQSFFSGLFGGSS KIEEACE\IYARAANMFKMAKN WSAAGNAFCQAAQLHLQLQSK HDAATCFV\DAGNAF\KKADPQ GGPFNCFDCVAFEIY\TDMGR\F TIAAKHHISIC*DLLRQKLVDIE KAIA\HYRALADYYQGEESNS\S ANKCLLKVAGYA\ALLEQYQK\ AIDIYEQGWGPNAHWTSPPPSK Y\SAKDYFF\KAALCHFCMDML QA\KLAF\QKYEELFPVFSDSRE CKLMKK\LLKA\HGEQNVD\SY TE\SVKGITDSISRL\DQWL\TTH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21968	52336	Α	22096	20	2482	RVCSSSASTASQAVMADAWEE
						IRRLAADFQRAQFAEATQRLSE
:						RNCIEIVNRLIAQKQLEVVHTL
						DGKEYITPAQISKEMRDELHVR
ļ						GGRVNIVDLQQVINVDLIHIEN
						RIGDIIKSEKHVQLVLGQL\IDEN
				-		YL\DRVGQEVNDKLQESGQVTI
ŀ						SELCKTYDLPGNFLTQALTQRL
						AIIIIGHIDLDNRGVIFTEAFVAR
						HRARIRGLFSAITRPTAVNSLIS
						KYGFQEQLLYSVLDELVNSGRL
						RGTVVGGRQDKAVFVPDIYSR
						TQSTWVDSFFRQNGYLEFDALS
				:		RLGIPDAVSYIKKRYKTTQLLFL
						KAACVGQGLVDQVEASVEEAI
						SSGTWVDIAPLLPTSLSVEDAAI
						LLQQVMRAFSKQASTVVFSDT
		}				VVVSEKF\INDCTELFRELMHQ
						KAEKEMKNNPVHLITEEDLKQI
						STLESVSTSKKDKKDERRRKAT
						EGSGSMRGGGGGNAREYKIKK
1		1	ļ			VKKKGRKDDDSDDESQSSHTG
						KKKPEISFMFQDEIEDFLRKHIQ
						DAPEEFISELAEYLIKPLNKTYL
						EVVRSVFMSSTTSASGTGRKRT
						IKDLQE\EVSNLYNNIRLFGKRG
						WKFF\ADDTQGWLLPKHLLEV
		1				QCVLISLNLI\FNF\LGFGILMDG
		1				Q*DDPGRPFTS\ELRKEI\LSKLS
						EETKVALTKLHNSLNEKSIEDFI
						FC\LDSAAEACDIMVKRGDKKR
						ERQILFQHRQALAEQLKVTEDP
21969	52337	A	22097	1	711	VLSLFDGIATGYLVLKELGIKV
	1		1207			GKYVASEVCEESIAVGTVKHEG
				İ		NIKYVNDVRNITKKNIEEWGPF
		ļ	ŀ			DLVIGGSPCNDLSNVNPARKGL
						YEGTGRLFFEFYHLLNYSRPKE
						GDDRPFFWMFENVVAMKVGD
]				KRDISRFLECNPVMIDAIKVSAA
	1					HRARYFWGNLPGMNRPVIASK
						NDKLELQDCLEYNRIA\RIFGFP
	1					VHYTDVSNMGRGARQKLLGRS
		İ				WSVPVIRHLFAPLKDYFAC
	<u> </u>	<u>. </u>	l	<u> </u>	<u>L</u>	WOVEVIRGILITATERDITAC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21970	52338	A	22098	406	2885	GRGILVNGACSDQSSDSPPILEA
						IRTPEIRGGWAVGTGVVSGADI
						VSGHCRQWSLLSLFQGSGTLSL
						AFITWAFRETRTRSESPAVSSHT
						SSQSTCGQHST*HT*HARGRQG
						RNHVDESPVEFPATRVGSPDAQ
						TPARSL*LGDMPHSLTIDLTDDT
						EDTHGTPQSSSTPYARLAQDSQ
						QGGMESPQVEADSGDGDSSEY
						Q\DGREGLLPRV/I*GP*TRGKTS
						YAMVVSWKATSKRQAMSGMR
						WVQWFGDGKFSEVSPGKGKGF
1						CRPEAVPACLPLNKLVSYRKA
						MYHALEVTWDEGMRAKP*EQ
						G*SKRPTELEDQLKPMLEWAH
						GGFKPTGIEGLKPNNTQPGGNE
						SPWQHPLPLLVGPLLGRVRLKT
						NCYNNGKDRGDEDQSRGDCW
						VPGIMGQMGGGRCRSGIDVIPL
						TPWLGL*ACLVCQTCRVSPPTT
						ALDLPPLASSNSPLLPCEGRELL
						LCSNTSCCR*ALGPVGWMWVG
	:				1	PQGSYVPAV*EPWSCYMCLPQ
					:	RCHGVLRRRKDWNVRLQAFFT
}			<u> </u>			SDTGLEYVSHRLPPLPPQIPGAQ
						GVGKLWFDGIATGEFGEHLETL
						LSCHNRVARESEESIAVGTVKH
			 			EGNIKYVNDVRNITKKNVRAV
						CTLRASSLPVFFPSPPHPNPARK
						GLYGEHPSLWQSLESLCHLTTG
						PGAGNK/RGLGWRFQMNPALS
						SFLST\GVQRDISRFLEVRESGD
21971	52339	Α	22099	729	896	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21972	52340	Α	22100	l i	2672	MAYKQGFLKVMVHFRKAEVT
						GKNHTSVHEGYTLISPAAQVLT
	1					ISCKKSILLQEQVSFKDVCVDFT
1		1				QEEWYLLDPAQKILYRDVILEN
į						YSNLVSVGYCITKP\EVIFKI/ES
			ľ			KGEEPW\VLEKGIPQAQCPPQK
						RKWKV/EMNVVREQPRKMKM
						TIFGEPPIPTTNKTVSVENGDRG
						SKTFNLGTDPVSLIKLSL*NM*L
		1				\SEMNLKNISGLIITKKNCSRKK
						PDEFNVCEKLLLDIRHEKIPIGE
	-					KSYKYDQKRNAINYHQDLSQP
	1					SFGQSFEYSKNGQGFHDEAAFF
						TNKRSQIGETVCKYNECGRTFI
						ESLKLNISQRPHLEMEPYGCSIC
						GKSFCMNLRFGHQRALTKDNP
						YEYNEYGEIFCDNSAFIIHQGAY
						TRKILREYKVSDKTWEKSALLK
						HQIVHMGGKSYDYNENGSNFS
1						KKSHLTQLRRAHTGEKTFECGE
į.						CGKTFWEKSNLTQHQRTHTGE
					}	KP\L*CTEC\GKAFCQKPHLTNH
						QRTHTGEKPYECKQCGKTFCV
						KSNLTEHQRTHTGEKPYECNA
						CGKSFCHRSALTVHQRTHTGE
		1				KPFICNECGKSFCVKSNLIVHQR
		1				THTGEKPYKCNECGKTFCEKSA
						LTKHQRTHTGEKPYECNACGK
						TFSQRSVLTKYQF\IHTRVKAL*
						TS*MLEAFIHL/SKLVIQFQKRRS
						EKAKECQKFVE\KTSCLNM*KL
						SRKIKTFH*KICKLTVKNKGNK
21973	52341	Α	22101	1	2817	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21974	52342	Α	22102	474	2766	NQTALYNQAQLLHGSRFANMS
					ļ	AEGYQYRALYDYKKEREEDID
	İ					LHLGDILTVNKGSLVALGFSDG
						QEARPEEIGWLNGYNETTGERG
	1					DFPGTYVEYIGRKKISPPTPKPR
	1					PPRPLPVAPGSSKTEADVEQQA
	İ					LTLPDLAEQFAPPDIAPPLLIKL
						VEAIEKKGLECSTLYRTQSSSNL
						AELRQLLDCDTPSVDLEMIDVH
	1					VLADAFKRYLLDLPNPVIPAAV
	Ì	İ				YSEMISLAPEVQSSEEYIQLLKK
						LIRSPSIPHQYWLTLQYLLKHFF
						KLSQTSSKNLLNARVLSEIFSPM
						LFRFSAASSDNTENLIKVIEILIS
	1					TEWNERQPAPALPPKPPKPTTV
						ANNGMNNNMSLQNAEWYWG
			İ			DISREEVNEKLRDS\ADGT\FLV
		1				RDASTKMHGDYTLTLRKGGNN
						KLI\KIFHRDGKYGFSDPLTFSSV
		1				VELINHYRNESLAQYNPKLDVK
	1					LLYPVSKYQQDQVVKEDNIEA
						VGKKLHEYNTQFQEKS\REYDR
						LY\EEYTRTSQEIQMKRPAIEAF
						NETIKIFE\EQCQTQERYSKGNY
	İ					HLEKFKP*RAMEKEIQRIMHNY
		ŀ				DKLKSRISEIIDSRRRLE\EDLKK
						Q\AAEYREIDKRMNSIKPDLIQL
		ŀ				RKTRDQYLMWLTQKGVRQKK
						LNEWL\GNENTEDHYSLV\EDD
						EDLPHHDE\KTWECLGSS/RTRN
						QSLKNLFARGSRDGTFLVRESS
						KQGCYACSVVVDGEVKHCVI\
21975	52343	Α	22103	3	392	
21976	52344	Α	22104	168	404	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21977	52345	Α	22105	49	1760	YQSPAWQSHWHQKLQVRKEN
						MAKRVAIVGAGVSGLASIKCCL
						EEGLEPTCFERSDDLGGLWRFT
						EHVEEGRASLYKSVVSNSCKE
						MSCYSDFPFPEDYPNYVPNSQF
		Ì				LEYLKMYANHFDLLKHIQFKT
	ļ					KVCSVTKCSDSAVSGQWEVVT
						MHEEKQESAIFDAVMVCTGFL
		Ì				TNPYLPLDSFPGINAFKGQYFHS
						RQYKHPDIFKDKRVLVIGMGNS
		l				GTDIAVEASHLAEKVFLSTTGG
						GWVISRIFDSGYPWDMVFMTR
					İ	FQNMLRNSLPTPIVTWLMERKI
						NNWLNHANYGLIPEDRTQLKE
						FVLNDELPGRIITGKVFIRPSIKE
						VKENSVIFNNTSKEEPIDIIVFAT
1						GYTFAFPFLDESVVKVEDGQA\
						SLYKYIFP\AHLQKPTL\AIIGLH/
						IKPLGLP*LPTGETQVRWAVSG
		ļ				S*KVLTRLPPPSVMIEEINARKE
1						NKPSWFGLCYCKALQSDYITYI
						DELLTYINTKPNL/FSLWLLTDP
						TSGLWTVFFGP\C\SPYQFRLTG
1						P\GKMGKGAQEIALHGPQW\DP
						NISRVHSKVRSCYKEFSIFPFGK
						FFLKVFSFPGFGLWLFFLIFPYK
21978	52346	Α	22106	262	969	EWSSVRRSLVEKRALRRPHPQC
						LCFRMKTIL\SNQTVDIPE\NVDI
						TLKGRTGIREGPPEGTLR\RDFN\
						HIN\VELSLLWKEKKRGSRVDK\
						WWGNQKRNWPTRSD*FVSHV
						QN\MIKGWLHWGFRLQG*GPV
						YAHFPHQPLLSRKNGVSWLKIR
						KFLWGEKYI\RRVSG*RPGCLL
						VSVSQGPRKDEINPLKG\NDIEL
						VSK/SSAGF*FQQATTS*KTRNI
						RKFL\DGIFCLLEKGLFRQA
21979	52347	В	22107	211	457	
21980	52348	Α	22108	1	159	KEAENQCLDISSPGMSGHRQGH
						DSLEHDELREIFNDLSSSSEDED
						ETQHQDEK*LDISSPGMSGHRQ
						GHDSLEHDELREIFNDLSSSSED
						EDETQHQDEK
21981	52349	С	22109	167	367	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21982	52350	A	22110	754	1950	VKGKDDAPHELGE/SQFILRLPP
						\EYASTVRRAVQSGHVNLKDRL
						T\IELHPDGRHGI\VRVDRVPLAS
						KLVD\LP\CVMEEPENPLLKKPF
						SKTAIIGQMLVSTVDGDLFPPV
						EEPSCLALDPKAS\KKKDKDKE
						KK\FIW\NHGITLPLKNVRKRRF
						RKTAKKEIYLNLPDV*KKELKR
						\LLEVPDA\EAVSTRWGIICPKV
						ETKG\AENQG\LDISSPGMSG\H .
						R\QGH\DSLEHDELREIFNDLSSS
						SEDEDETQHQDEEDINIIDTEED
•						LERQLQDKLNESDEQHQENEG
1						TNQLVMGIQKQIDNMKGKL\QE
		ŀ				TQDRAK\RQEDLIM\KVGNLGS
						PRTDFQAVLGMSFNQREDREK
						EQL\SSLQEELESLLREVKKNLM
						FNFQSFRLVQHLGKFFGLLLLL
						GYLRPCSS
21983	52351	В	22111	227	355	
21984	52352	Α	22112	121	486	WGPALCIGGYLASFLASVHLMP
				1		LR*GEMLSP*GLSFSFRFPGFKE
						VRLVPGRHDIAFVEFDNEVQAG
						AARDALQGFKITLSPLKVSPPL
						GAFLEPCVSEWSPHSIVPRVCPQ
		<u> </u>				TLHLALLGRN
21985	52353	В	22113	8	165	
21986	52354	Α	22114	1	218	
21987	52355	Α	22115	566	1412	TLHSMAVPETRPNHTIYIN\NLN
						EKIKKDELKKSLLRPSFF/SSLG
						QILGYSWGIHGRP*RMEGARPF
						SIFKEVSSGTNALR\SMQGFPFY\
						DKPMRIPVLPRTRLQDIHLPRIE
						RAPFIGSRDPKAGERGRPKSQG
						DPGPPKEGLCKGGGSHPRGGG
						LSRGPGPGHCPPMT\KAPRI\MH
			1			HMPG\QPPLTAAPLDMIPPPG\L
			1			APGQIPPG\AMPP\QQLMPGQMP
]			P\AQ\PLWRNPP\NHILF\LTNLP\
]			EET\NELMLSHAFSNQVPPGFKE
						GPVWVPRAGNEHRPSVEV

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21988	52356	A	22116	2	1176	FGELSTFDAVRGPPRRCPPDAD PRAPGLALPGRLLVSNSRRTHS RESRAPPAPHLLGARAGMAPPQ VLA\FGLLLAAATATFAAAQEE CVCENYK\LAVNCFCE*INRQC Q\CTSVGAQNTVICSKLGCQMF W*MKAEMNGSKLGRRAKPEG GLQ\NNDGLYDPDCDESGLFKA KQCNG\TSTC\WCVNT\AGVRRT HKDTEVTCSERVRTYWIIIELKH KAREKPY\DSKS\LRT*LQKEITT RYTLGSKFLSRVFWYENNVITI\ DLFQNFFSKKLRNDVDIADVAY YF\EKDVK\GESLFH\SKKMDLT VNGEQLDLDLLIPGQTL\IYNVD EKAP*/DFSMQGLK\AGVIAVIV VVVITSCCWKLLVLVISQKGRE WQKYEKAEIKGDGCRCHRELQ
21989	52357	A	22117	3	257	AKALQWPLGRGGGGTPRGSRP PGPH*SERGAVQPRSLPRQAA* RVPSGPADGQ*DGHGAADPGS RQRCKPWSEAADLHSECYSS
21990	52358	A	22118	1467	1748	SEYNCCEIGCPQPSWGPWGRG\ TDVRQGQVGEEEVHG/ECGGG SQG*RPG**AGSPAP*PGTWTGT AQQGPAAVLDPLRV*GGGIERH PLHSVGL
21991	52359	В	22119	1	1461	
21992	52360	Α	22120	29	417	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21993	52361	A	22121	616	1671	TATCKPWSMRTTISSSQPQKLT NSVKLPPSAKCISVSRIGGFLVS LTSRMKLRTLLICSKLLPSGGV RGLAGSGVKLQTFTVSVTALK AACLELFTSPGGFMVLLASEVK LQTFSAGDSDCLGQGRRAQRRP GPGVSAVGASGQQRAPESNERP AEEPTPAPMPSSAPRGPTEPALR RPSSADRPLAPGAGRAPGAMAS PSGSSKATGKPRGRDGRPRREE DDVTPEEKRLRLLLEEGSAQPD DGEDAPRPGRKETGTQTGGDG RGAARAALLAVGGIRAPLPSLG VAPGC\PMSKPPRRHPGATPR/P GQGGSDAAHGEEGCPCCSSAV ATCLGAGLLP\PGRGASSPSSGC ALPSSSNSRSLFSSGVTSSSSLLG RPSLPRGLPVALELPEGLAMAP GALPAPGASGLSAELGRLSAGS VGPRGA\G*ASGRAWAPPRAAR CSLAPSAG\PRAHRGHAGPWPA LSSP/GPGPSSRCPLPRRSAASLL KPAR/AINPPGEVNNSRHAALR AVTLTVKVCSFTPEPARPRTPPE GRSFEHIRRVRSFILEVSETKNP PILDTLIHLALGGSFTLFVNFCG
21994	52362	A	22122	892	1931	WDELIVVLIDQGLHVAV CTWNENQRMEIVWEVLFLLQA NFIVCISAQQNSPKIHEGWWAY KEVVQG\NFVPVPSFWGLVNSA WNLCSVGKRQSPVNIETSHMIF DPFLTPLRINTGGRKVSGTMYN TGRHVSLRLDKEHLVNISGGPM TYSHRLEEIRLHFGSEDSQGSEH LLNGQAFSGEVQLIHYNHELYT NVTEAAKSPNGLVVVSIFIKVS DSSNPFLNRML\NRDTF\TRN*H FKNDGIFYYRGL*YRGELLFQG TSSFITYDGVDGLFPTPGY/SRT ASWIIMNKPV*YNPGCKMHSLR LL\SQEPAISRSFLSMSDNFRPV QPLNNRCIRTNINFSLQGKDCP NNRAQKLQYRVNEWLLK
21995	52363	A	22123	17	417	THE TALL OF THE TELES
21996	52364	A	22124	63	1131	CHPALLGPTGHGAWPRHRAPD TLRASAGGRDP*GRRSAPA*ER QP*PPVYCAGISQYD*TALSSCL RRSVRTSTDFRSLLDKSNYSVF LCFLQLAGRRYFAMWTRPEFFL ARDPRSLSWGLEWRPLSL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21998	52366	Α	22126	359	548	
21999	52367	A	22127	755	1031	TRGVLEHMALGKGTLKDPSGP AP*NHPE\VSHAASTQEAAGPP AGRPGHGSGCLALPASSTSLEP GLRQDHDLLLQLLWREQEVFH DVIQLL
22000	52368	A	22128	1	277	
22001	52369	Α	22129	1	933	
22002	52370	A	22130	339	863	PSPVPARPVCPSSAPISMRPPCP WREAEAADLRPEAPGHRDGED VGFLSSGQLSGLCSKRKLQPTR AALTLTPSAVNKIK\HFLKNNPE HVGVKVGVRTRGCNGLSYTLE YTKTKGDSDEEVIQDGVRVFIE K\KAQLTLL*TEMDYVEDKLSK GVGV\ITPNIKGTCGCGESFNI
22003	52371	A	22131	16	419	QIQHRPWP/SRSYTNVLGRKVV YLPSFFTYAKYIVQVDGKIGLF RGLSPLMMSNALYTATRCFMK KDLPPDEIEQVSNKDDMTTSLN KVANDTSYEMMMHCVSRMLP HPLHVISMRCMDQIVGREAKYS GVLSSIG
22004	52372	A	22132	3	441	RSYTKFVMGIAVSMLTYPFLLV GDLMAVNNCGLQAGLPPYSPV FKSWIHCWKYLSVQ\GASTGRR RPFLFFATSFS*NGFVDASLHAK GQLFRGSS\CFSAGCHQDHALP WSNLNHLKNTVSTWPPWVRPD HLGTPARRLQPNNNQM
22005	52373	Α	22133		1232	MAGAGAGAGARGGAAAGVEA RARDPPPAHRAHPRHPRPAAQP SARRMDGGSGGLGSGDNAPTT EALFVALGAGVTALSHPLLYV KLLIQVGHEPMPPTLGTNVLGR KVLYLPSFFTYAKYIGQRDG\RE GLVPRLSPRLMSNALSTVTRGS MKKVFPPDEIEQVSNKDDYEDF P*RKLLKETSYEMMMQCVSRM L\AHPLHVISMRCMVQFVGREA KYSGVLSSIGKIFKEEGLLGFFV GLIPHLLGDVVFLWGCNLLAHF INAYLVDDSVSDTPGGIGKRPE SRFP\FSQALALRSYTKFVMGIA VSMLTYPFLLVGDLMAVNNCG LQAGVPP\YSPVFKSWIHCWKY LSVQGQ\LFRGSS\CFSAGCHQD HALPW\SNLNHLKNTV\STWPP WV\RPDHLGTTARRFQPNNNQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22006	52374	A	22134		1156	PRHPRPAAQPSARRMDGGSGG LGSGDNAPTTEALFVALGAGV TALSHPLLYVKLLIQVGHEPMP PTLGTNVLGRKVLYLPSFFTYA KYIVQVDGKIGLFRGLSPRLMS NALSTVTRGSMKKVFPPDEIEQ VSNKDDMKTSLKKVVKETSYE MMMQCVSRMLAHPLHVISMR CMVQFVGREAKYSGVLSSIGKI FKE\EGLLGFFVGL\IPHLLG\DV VFLWG\C\NLLAHF\INAYLVDD SFS\QAL\AIRSYTKFVMGIAVS MLTYPFLLVGDLMAVNNCGAA SWGSPPYSPVFKSW\IHCWKYL SVQGQLFRGSS\CFSAGC\HQDH GFALGVN*NHLKNT\VSTWPPW VRPDHLGTPARRLQPNNNQMC SSPAG\FQFHICHVSV\QMWG
22007	52375	A	22135	2	425	SSI NOSI QI INCII V SV QII W G
22008	52376	A	22136	2	430	
22009	52377	A	22137	2	406	
22010	52378	A	22138	1	297	
22011	52379	A	22139	1	111	
22012	52380	A	22140	1	398	
22013	52381	A	22141	1	409	
22014	52382	Α	22142	500	1234	GPNCRPPSCSAVEGAWPRPQGA GARAAIALSQKLELGGVGLVQ VQCPSLAPSSTSSILFLCLLPHPP PHRCLQENTTCLLGVEGVLLLQ SPFPLLKTDHSSLPALAPTQSQA ALWGFADFGFMFESSPGGNLG WEPDIKLTDEMVMIMGGKMEA TPFKWFMEMCVRGYLAVRPY MDVVVSLVTIMLDTGLPCFRG/ QIKFLKHRFSPNMTEREA*NFIM KVIQSCFLSNRSRTYNMIQYYQ NDIPY
22015	52383	A	22143	2	6378	SAGPAGPLQVQKLLCMCPVDF HGIFQLDERRRDAVIALGIFLIES DLQHKDCVGPYLLRLLKGLPK VYWVEESTARKGRGALPVAES FSFCLVTLLSDVAYRDPSLRDEI LEVLLQVLHVLLGMCQALEIQ DKEYLCKYAIPCLIGISRAFGRY SNMEESLLSKLFPKIPPHSLRVL EELEGVRRRSFNDFRSILPSNLL TVCQEGTLKRKTSSVSSISQVSP ERGMPPPSSPGGSAFHYFEASC LPDGTALEPE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ľ		sequence		
22016	52384	Α	22144	3	398	 IHKAASGVPQQPDAGCPGTGEQ
						QGGQRCGQ*AGP/AQAPC*SRA
						GTR/PTCFFVLTSLRSWYRVDSP
						NKLMNPLVAGFFG/CHCGSGQC
						LRKCSSARDRAPDARPGLDLAQ
		ļ				GRCCHCNKACLHASETGVGTT
						SVL
22017	52385	Α	22145	344	498	
22018	52386	Α	22146	344	406	FLADQEDPLDPRAGCR*ETTAP
ŀ						RIPSKGPSSTEPLHASTAVPPCQ
İ						GELGCSRASPASEKI/DLAGRLIS
						PCNSDTVSPGLGGFMTCAQSLI
						SRRPGGPIGPPGWLQGPHLG
22019	52387	Α	22147	3	2058	LSGSGAPPPGVGAQAAAAAEE
						EEREVVRVRVKVRLAVSRPSRR
						GKFRSFAVDPQITSLDVLQHILI
į						RAFDLSG*VGRCWGLLGRDRL
ŀ		1				GQEVYLSLLSDWDLSTAFATAS
ļ						KPYLQLRVDIRPSEDSEYSVPPG
į						AISPKDVIGSDVLLAEKRSSLTT
						AALPFTQSILTQVFSGTLGTQQ
		1				VLSWSYGEDVKPFKPPLSDAEF
İ						HTYLNHEGQLSRPEELRLRIYH
						GGVEPSLRKVSILSHLLVYPDG
		1				LTGRERMDYMKRKSREYEQLK
						SEWAQRANPEDLEFIRSTVLKD
						VLRTDRAHPYYAGPEDGPHLR
						ALHDLLTTYAVTHPQVSYCQG
						MSDLASPILAVMDHEGHAFVC
		l	1			FCGIMKRLAANFHPDGRAMAT
						KFAHLKLLLRHA\DPDFYQYLQ
						EAGADDLFFCYRWLLLELKREF
						AFDDALRMLEVTWSSLPPDPPE
					}	HEVELVGPPSQVADTGFGDHR
						G\WPVRQRHMLRPAGGGGSTF
						EDAVDHLATASQGPGGGRLL
						RQASLDGLQQLRDNMGSRRDP
						LVQLPHPAALISSK\GLSEPLLN
		1				YPDPLLSSFSHPDSPSSSSPPSTQ
		1				EASPTGDMAVGSPLMQEVGSP
						KDPGKSLPPVPPMGLPPPQEFG
						RGNPFMLFLCLAILLEHRDHIM
				ĺ		RNGLDYNELAMHFDRLVRKHH
22020	52388	<u> </u>	22148	130	207	LGRVLRRARALFADYLQSEVW
22020	52389	A A	22148	1	387 279	
22021	32309	Λ	22177	<u> </u>	217	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22022	52390	A	22150	105	499	DMSDTSESGAGLTRFQAEASEK
22022	22370		22.30	""		DSSSMMQTLLTV/TPEAREAPA
						TQASSTTQLTDTQVLAAENKSL
						AADTKKQNADPQAVTMPATET
		İ				KKVSHVADTKVNTKAQETEAA
						PSQAPADEPEPESAAAQSQENQ
						DTR
22023	52391	Α	22151	285	984	TSLSCYFSRTKDSPKLGLLMVL
						LSIIFMNGNRSSEAVIWEVLRKL
	•					GLRPGYDWALSALAVRVVLW
						QERTVLGLHQSGGLVERVGCW
1						TG*RAQGSDLGG*WVNGPELS
						APSLSVSWASMELPSCAGNLFS
						ILEMPLPTSGKCHSLE*IYLFIYF
		1				SFSSLRIHHSLFGDVKKLITDEF
						VKQKYLDYARVPNSNPPEYEFF
		İ				WGLRSYYETSKMKVLKFGCKV
						QK\KDPKEWAAQYRE
22024	52392	A	22152	259	578	
22025	52393	Α	22153	296	2242	NMSDTSESG\AGLTRFQA\EASE
					•	KDSSSMMQTLLTVTQNVEVPE
						TPKASKAL\EVSEDVKVSKASG
						VSKATEVSKTPEAREAPATQAS
						SYLLSLTD\TQVLAA\ENKSLAA
		l				*HQGNRMVDPQAC*QLPAT\ET
						KKVSHVADYGRFNYKGFRRLR
						LAPS\QAPADEPEPESAAAQFSG
						RFRVFGPKVKAKKARKVKHLD
ĺ						GEKDGSSDQSQASGTTGG\RRV
į						SKALMASMARRASRGPIAFWA
	1				<u> </u>	RRASRTRLAA\WARRAFALPLEI
	1					TLKPRRAKGLRR\RVAKFQ\SSQ
						EPEAPPPRG\LALLQGRAN\DLV
			ŀ			KYLLAKDQTKIPIKRSDMLKDII
	1					KEYTDVYPEII\ERAGYSLE\KVF
						GIQLKEIDKNDHLYILLSTLEPT
						DAGILGTTKDSPKLGLLMVLLS
						IIF\MNGNRS\SEAVIWE\VLRKL\
	1					GLRP\GIHHS\LLGDVK\KLITDE
						F\VKQKYLGLCQKVPNSNPP\EY
						EFLLGACALY**EPAKMKSPQS
						FACK/VYQKEGSPRNWGSSVPP
						REGKGRMEADFEGCKLRLTLE
						AKAKGPRLEAPNGPLGLGSGE
						MLAGP\CNWDEA\DIGNPWAK
						ARIPRREQKLKAKAPKRSGSA\S
						TGA\STST\NNSAS\ASA\STSGGF
						\SAAASL\TATLHIWGSSAGLWL
						ELGCQHQSARSGAC\GVSYK
			•			· —

SEQ ID		1	SEQ ID NO:	1	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deterion, '-possible nucleotide insertion)
				sequence		
22026	52394	Α	22154	1	487	EFRQPSFRRSIWP/YFLLTQA/VR
						TGNPAKFNQVLDQFGEKFQAD
						GTYTLIIRLRHNVIKTGVRMISL
						SYSRISLADIAQKLQLDSPEDAE
						FIVAKAIRDGVIEASINHEKGYV
						QSKEMIDIYSTREPQLAFHQEG
						KDSSVYSFREWGTRTGMSWLV
		_				CFLKLRL
22027	52395	A	22155	2	429	
22028	52396	Α	22156	1	328	GTRFQADGTYTLIIRLRHNVIKT
						GV/RSGSEGLLEEQKPGQGKPS
	i					GYPWASEGSVSTLPTPSLLPLPG
						VRMISLSYSRISLADIAQKLQLD
						SPEDADFIVAKVGLVQEPQ
22029	52397	Α	22157	138	2206	AVTPGAMKQEGSARRRGADK
						AKPPPGGGEQEPP\PAPAPQDVE
				1		MKEEAATGGGSTGEADG\KTA
						AAAA*SNFPAKSWDYSHLWKD
						IKEHRETS*RKAVSGKEPRFVLR
				1		ALRMLPSTSRRLNHYVLYKAV/
				}		QGAFFTSNNAT\RDFLLPFLERA
						HGTQEADLLFRPRTGKAAGT\P
						LLPEVEAYLQLLVVIFMMNSKR
						YKEAQKISDDLMQKISTQNRRA
						L\DLV\AAKCYYYHARVYEFL\D
						KL\DVVRISLHG\RLRTATLRHD
						ADGQAT\LLNLLLRNYL\HYKL
						VPTSLRSLVSKSVFP/EEQAQQP
						NEWGQVTSYLHQGRIKAIQ\LE
						YSEARRTMTNALRKAP\QHTAV
						GFKQTVHKLLIVVELLLGEIPDR
						LQFRQPSLKRSLMPYFLLTQAV
						RTGN\LPRFTQVLDQFGEKFQA
						DGTYTLIIRLRHNVIKTGVRMIS
						LSYSRI\SLADIAQKL\QLDSPE\D
						AEFI\VAKAIR\DGVIEASIN\HEK
						AYVQ\SKEMIDIYST\RDPQLAL
						H\QRISFCLDIHNMVWSRPMRV
						PSPKSYKQGSWSLQRNGRERR
						NKQDLE\FPKEMA\EDDDDSFPL
						KLGGWGGVRGKWGTRQLCPP
						WGYPLPQGTCPPFSPHTAHML
				1		AFRGRGWGCWEPATPDLPPGL
						LPSRVTLLVQPGRRVGRQPPRG
						RVLASSVADRRGRISLCTPFRE
		<u> </u>				WGTRTWDVYGLLCFLKLRL
22030	52398	Α	22158	86	348	

SEQ ID NO: 22031	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion) LPLSMDVRAPPAVMAAVDQAL
22031	32399	A		3		KEFGRIDILINCSSSSCGLPFCR\ AAGNFLCPAGALSFNAFKTVM DIDTSGTFNVSRVLYEKFFRDH GGVIINITATLGNRGQALQVHA G\SAKAAVD
22032	52400	A	22160	3	255	GPWSCHAGA*GPGATQVSEDQ GLREPHGSRDLCSLG*KH*GAPI CAWPMSWDGPPAAHKVRPSVS APPVCINTLCYNLASAFS
22033	52401	Α	22161	3	360	
22034	52402	A	22162	240	1356	AGNPSGIPGTRVHRGQRGGFAS TWSGRGAGSTPLQNSPPRQPHR EKEFLRQVLEGGSHTGVFRPSA ARLVPNPHSLVSPDLL*LVSRA T\PHIFLAARKLAGATGRRCLPL SMDVRAPPAVMAAVDQALKEF GRIDILINCESVLSEVGGFSQAP AAPAVSHSAGAAGNFLCPAGA LSFNAFKTVMDIDTSGTFNVSR VLYEKFFRDHGGVIVNITATLG NRGQALQVHAGSAKAAVDAM TRHLAVEWGPQNIRVNSLAPGP ISGTEGLRRLDASQGMSWGRG EGAGSWGSRGLSLLLPS\GGPQ ASLSTKVTASPLQRLGNKTEIA HSVLYLASPLASYVTGAVLVA DGGAWLTFPNGVKGLPDFASF
22035	52403	В	22163	214	1647	

2846

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22036	52404	A	22164	1765	1910	LSHPHSRTPPNRTPRRTRIPQRP
						AVMYSPLCLTQDEFHPFIEALLP
						HVRAFAYTWFNLRARKRKYFK
						KHEKRMSKEEERAVKDELLSE
						KPEVKQKWASRLLAKLRKDIR
						PEYREDFVLTVTGKKPPCCVLS
						NPDQKGKMRRIDCLRQADKV
						WRLDLVMVILFKGIP\LESTDGE
						RLVKSPQCSNPGLCVQPHHIGV
		i				SVKELDLYL\AYFVHAGRISSQS
						ESP\SQ\PSDADIKDQPENGTFGA
						SRDSFVT\SGVF\SVT*A*LRVSQ
ŀ						TPI\AAG\TGPNFSLSDLESSSYY
						SMSPGAMRRSLPSTSSTSSTKRL
						KSVEDEMDSPG\EEPFYTGQAR
						SPGKWTVSPSGLSVSWSQGMPS
						PTTLKKSEKSGFQQPLPFHRPPS
		l				LGNGFHTAFTDLFIYQGPQSKF
						HIATPSIL\HFPRHSPFFQQPGPY
		ł				FSHPAIRYHPQETLKEFVQLVCP
						DAGQQAGQPNGSSQGKVHNPF
						LPTPMLPPPPPPPMARPVPLPVP
		1				DTKPPTTSTEGGAASPTSPTTRS
		1				/PGRTRPQQPFL/SYGPP*PHFLR
						AGIHHL
22037	52405	Α	22165	63	250	
22038	52406	Α	22166	1	817	SRLGHPRKDMDDDTKATLISY
						MSLSFRHGYCFSEDPCVHTELG
						IQSGRGGRGDPDPPGNALE*LA
		ł				HLGP/SSHLGP*AQDGLVDAAG
						RASPLAGTEGAGRFSHLQPI/GP
						RGPPE*GAAGERFPSGVAG/GPR
						GAHGLPCPEQLQLQPPPPS*MP
						KTGSTFVAHFPSPLAQGHSPPG
		1				GGAGWCSGCSPDRPPLLGVGS
						HGELLNPNSPAYEVGAG/RPAT
Ì		1		}		GSSGTRAGPPSPSHSAPGRTPSV
						SLCD/RHQCLRASPTGSIGSLPA
		L				HWREGLAGPLMD

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	""	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22039	52407	A	22167	240	1790	TAKNSYLESCPLHMENSIHQWP
:						RQPFPPVRRQGADAACGAGSQ
						ALVPSHSDTDGVLPGAEWLGD
						GGPARVP/GAAGGRGPAPTS*A
						GP*LSL*APLCF\SGLSGNGKQL
						LGAAGRSSSHAQMHELHPAQ*
						HE*ATDHTGGQGPP/G*KQQNG
						PP/GPDNDLPAPESTPPTWNEDF
						LPDAIPLAHPVPRQRRPTGPAGP
						PGQTGPPGPAGPPGSKGDRGQT
						GEKGPAGPPGLLGPPGPRGLPG
						EMGRPGPPGPPGPAGNPGPSPN
						SPQGALYSLQPPTDKDNG\TQG
						WPLPSWTQCW/SGVPGPRGPPG
	ļ					PPGPPGPRGPPGPPGTPGSQGLA
						GERGTVGPSG/AGGAAAERGPE
						DPGRASPHPGAHDWDPRSPGLP
						RGRFWPGCCPESQPQDEEGWH
						PTRWGPCCPAWARPWTEERG/S
						RPAAGS/RGVSDISETPGRPIYPD
		:				DKSQSAVSVLCARSRLRAQLRP
	ŀ	İ				GAEQMNDCKGITGTRQDIHAV
		i				SFASEHNIQNWVSKHLQLSPPV
						TPATQAPLKVRVPQASTLVL
22040	52408	Α	22168	104	271	PSCTNPYGGGRRCDSVVPNQCF
						TKHMQSRLLG*GEKALEGGGG
						RGVKSLVTKRDF
22041	52409	Α	22169	494	1074	KGTTLCQKHWGGGYLHWG\HF
						EMMRLTIN\RSMDPKNMFAIWR
						VPSPFQ\PITRKSVGHRM\GGGK\
						GAI*HYLTP\VKAGRLV\VEMG
						GRCEFARSAKVSLTQVALRSCP
						FASKRL*ACGTPREGCEQDQEE
						RERQQPRTPWTI*AE*PTAQHG
						WAYGKVPEPHMTLTPQGEILG
						ARFYMPKTCVVSVGDILYIGY
22042	52410	Α	22170	61	329	NQRSGKKSQPW*STITPPSLIWL
						SNSSTTLSSSKMGQSYSSS*PSY
						NCSVSRMFGSNMVPAATASGA
						SLTCPLLWLSMTHFRRKVGGW
						Α
22043	52411	Α	22171	21	444	STAQSISRSDLHYFLALLVYYW
						AADARFLPGRIPWLIHKELEKS
						HTEYCLVGSSQYTPLVQVFRSN
						TLVSVLSTKTI*NQRSGKKSQP
						W*STITPPSLIWLSNSSTTLSSSK
						MGQSYSSS*PSYNCSSRAKSPTL
						TVFQS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22044	52412	A	22172	163	1056	LEKSDHRISWSPGKTA\SPKSMP ERWHR*WAQILKD\MGIT\EYEP R\VIN\QMLEF\AF\RYVTTIL\DD AKIYSSHAK\KATVD\ADDVPIG QSRC\RAGSSLFTSPPPKRF/LLL RYLARQKKSNPFCPLIKPY\SGP RLPPDRYC\LTAP\NYRLKSFTE KGHSTSAGKELTVP\RLSVGFRL LSRTKVLPTLGAHPTPQ\TMSVF NLK*GTPHVPSPGQRFTSTRCPT S\QS\PAVKSFQFPGNLSQFQNV L\INPIH*IGSQKHSYYPLNMMS S\QNTA\NESSNALKRKREDDD DG\DDDDDDDDDDDNL
22045	52413	Α	22173	215	344	
22046	52414	A	22174	302	1330	ACEQKRGGEGGNKNRERRKGE KAGRRQGTRQPCSRCMRRTKQ VEKNDDDQKIEQDGIKPEDKA HKAATKIQASFRGHITRKKLKG EKKDDVQA\AEAEA\NKKDEAP /VFADGVEKKGEG\TTTARSSPQ PLAPKP\DEPGKAGETS\SEEKM GEG*LLATEQGSPPRLLAS\SEE KAGSAETESATKASTDNSPSSK AEDAPAK\EEPKQADVPA/AFG VFPNPLGALLLAATTPAAEDAA AKATAQPPTETGESSQAEENIE AVDE\TKPKESARQDEGKEEEP EA\DQEHAWNSGEMGFPHPHPP GCLACLSLPSSQLHSEVPSCPAH VCESVLSHPLALFLSVWANI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
<u>.</u>				sequence		
22047	52415	Α	22175	2	1577	HGFRLPHPQPALRRLLDQRRPK
				1		TEDRCGHWDQILDLSIIKHLFS
	}					MSFIPVAEDSDFPIHNLPYGVFS
						TRGDPRPRIGVAIGDQILDLSIIK
:						HLFTGPVLSKHQDVFNQPTLNS
•						FMS\LGHAAWKEARVFLQNLL/
ŀ						SL*SQARLRDDTELRKCAFISQA
i						SATMHLPATIGDYTDFYSSRQH
		ł				ATNVGIMFRDKENALMPNWLH
ŀ						LPVG\YHGRASSVVVSGTPIRRP
		}				MGQMKPDDSKPPVYGACKLLD
ŀ	1					MELEMAFFVGPGNRLGEPIPISK
						AHEHIFGMVLMNDWSARDIQK
						WEYVPLGPFLGEEFWGPLSLP
	}					WGGCPWDAL\MPF\AVPNPKAV
				;		PQAPCRYLCHDEPYTFDINLSV
				}		N\LKG\EGM\SQAAT\ICKVQILK
•		l				YMYWT\MLQQAHSPTLSNGC\N
		1				LRPGGPPGLSGTHQAGPEP/EKN
						FGSMVE\LSWKGTKPI\DLG\NG
						QT\RKFLLD\GDEVIITAMATRD
	1			İ		FKWYC\QGDG\YRIGFGQ\CAGK
						VLPALLPIMRFSLLFWKQRAQS
		<u> </u>				TPFQPCDWGSSPRAVGLVPPFQ
22048	52416	Α	22176	3	623	DSRNSSILPRRGALLKVNQELA
						GYTGGDVSFIKEDFELQLNKQL
1					1	IFDSVFSASFWGGMLVPIGDKP
						SSIADRFYLGGPTSIRGFSMHSI
				•		GPQSEGDYLGGEAYWAGGLHL
						YTPLPFRPGQGGFGELFRTHFFL
						NAGNLCNLNYGEGPKAHYIVK
		1				LAECIRWSYGAGIV\LRLGNIAR
						L\ELNYCVPMGVQTGDRICDGV
						QFGAGIRFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
22049	52417	IA	22177	352	2397	GPMSEGIASQGARRRRGIMGLR
22049	32117	 	22177	332	2337	GLWASRADGRRGETEPEAKQEI
						LENKDVSFLLKVLIELLVETFTS
						IHH*GDDVFKAKNLIEVGVVST
		İ				WCAFPVSSEELVNVPM*NIDTC
Ì		l				QGTYCGVVSL*FPFCPSEKSCFY
		l				GONNTMVGNNEGSMVCYRLF
						TFLYWNLELGTFTL*LHSKETS
						YGLSFFKPRPGNFERK*EAOOII
						EYTGLIEKLK*VAYL*KSAEYS
	;					VS\AFQSFPLGLGENRVAFSLIP
						KGSYLIQ*VSNGCWHLHCQPLL
						WGSYHRILAELRVLIESTWPFE
						G*LCNGTSQAFSCGPCTPSRSSA
						RHGHSVCGQLLVADVPGVKG
						MLGENNHLFLKY*SMNG*FA/L
						YFPLDLFSCFL*ESFPLTAPSLEG
						SRLTLETMYLLCQGGFGELFRT
						HFFLNAGNLCNLNYGKTCAIQE
						TIVVQLFSCLKAQYTKIRLGNIA
						RLELNYCVPMGVQTGDRYVLG
						IIFHNHIPLSSNFILFWTWFASGD
						TAGAARLLRLPTDDRAFRDGG
1						GCSHNRQLPFPSTPLLFSTQSPS
						WKGCSHAGCMHVCLGAQLPG
						LGAQLGHGSESQVERLLACGG
						RWGARRQGCCWHDLPKAPHA
						GCFLAGEKKVGDDQEPCTKTA
						AVREGESVGCTAEGGCLGLSG
						GCQGWLAPGLVELRSAAARQE
						GSRRRKRRRHLLRVHPQKVV
22050	52418	A	22178	1	1017	
22051	52419	A	22179	1	435	
22052	52420	Α	22180	1	1353	
22053	52421	A	22181	1	834	
22054	52422	A	22182	2	2675	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	, ,
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	:			sequence		
22055	52423	Α	22183	72	1339	AMRGANAWAPLCLLLAAATQ
						LSRQQSPERPVFTCGGILTGESG
						FIGSEGFPGV\WPPESKCTWKIT
						VPEGKVVVLNFRFIDLESDNLC
						RYDFVDVYNGHANGQRIGRFC
						GTFRPGALVSSGNKMMVQMIS
						DANTAGNGFMAMFSAAEPNER
						GDQYCGGLLDRPSGSFKTPN\W
						PDR\DYPAGVTCVWHIVAPKNQ
				1		LIELKFEKFDVERDNYCRYDYV
						AVFNGGEVNDARRIGKYCGDS
						PPAPIVSERNELLIQFLSDLSLTA
						DGFIGHYIFRPKKL\PTT\TEQPV\
						TTTFPVTTGLKPTVALCQQKCR
						RTGTLEGNYCSSDF\VLAGTVIT
						TITRDGSLHATVSIINIYKEGNL
						AIQQAGKNMSARLTVVCKQCP
						LLR\RGLNYIIMGPSRVKDGAR
						QNSWPNSFIMMFQT\RNQKLLG
,						CP*KISQC
22056	52424	Α	22184	49	170	HTVGDGRLLRRP*REFGASFL*
						CFLEARFGLKNWGHCWRH
22057	52425	В	22185	597	651	
22058	52426	Α	22186	1	791	
22059	52427	Α	22187	2	943	TGACAEPRVGLLFRLKGRCRG
						GRNMELGSCL\YGFREAAEEEG
						EPEVKKRRLLCVEFASVASCDA
				1		AV\AQCFLAENDWEMERALNS
						YFEPPAEES\ALERRPETISEPKT
						YVDLTNEETTDSTTSKISPSEDT
						QQENG\SMFSL\ITWEYLMGL\D
						LKQSVQRRARGVCSYLAL*VLS
						LLFNSNVSYVYTERMNQLKMV
						LKKMQEAPESATVIFAGDTNLR
						DREVSDKVSLMFEDRLLLWRLI
						*VPFCQYTWDTQMNSNLGITA
						ACKLRFDRIFFRAAAEEGHIIPR
						SLDLLGLEKLDCGRFPSDHWGL
	<u> </u>	<u> </u>				LCNLDIIL
22060	52428	Α	22188	277	534	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22061	52429	A	22189	29	1305	VGVVLRVECPLCPGPAPMSVPA
						FIDI\SEEDQAAELRAYLKSKGA\
ŀ						EISEENSEGGLHVDLAQIIEACD
	İ					VCLKEDDKD\VESVMNSVGIPY
						SWIL\EPDKQ\EAFDLKALCEKA
						GSKFPRR*TAPSLETCSFLSNLF
						H\GMG*GILPVKIHSVICSL\IKV\
						AASCG\AIQYHPQLEPGIKFRK
İ						WDFLNWN\LTTGKKSTPLLKTT
						FMKALVE\CKKSDAASKGM\VE
						LLGSYP\EGNASQ\ARVDAHR\C
						IV\RALKDPN\AFLFD\HLLT\LKP
					į	VKFFGKASLFHCSFLTKF\VSA*
1						MGHHMSRFYPD*LRTFH*IPLG
						PVTMDPGIWPKMRLLF/TFMGN
1						GQ*KNKEISF*PQCQQEPSRLGA
1						D\DVESHLVIDA\VRN*KWVYC
						QNWSRPRRKVVVSHSTHR\TFG
						K\QQWQTTVLTHLNAW\KQNL
22262	50.100	ļ.,	22122	104	1410	NK\VKNSLLSLF
22062	52430	Α_	22190	194	413	EVALUATION OF CACACA PACCACA
22063	52431	Α	22191	3	160	EVKFQWVLSLG*SFSLPKSSFQ
						RCGAGGAVHWSTALFLLGAAS
22064	52432	-	22102	268	473	ASSYHQRPG
22064	52432	A C	22192 22193	93	500	
22065	52434	A	22193	2	398	FFLRRSLALSPRPDCGLQWRNL
22000	32434	A	22194	2	370	GSLQAPLPGFTPFSCLSLPSSWD
						YRRPPPRPANFLYF**RRGFTLL
1						ARMVSIS*PHDPPASASQSAGIT
						GVSHRARPETYNFCSFIIISSLPV
						EPEARKMTKNSDIGEIW
22067	52435	A	22195	972	2180	PRHCCASASSAPAPSRPASSSSP
22067	32433	A	22193	1972	2100	PAACPW*GSLLGPAATPATWR
						R GSLLGFAATFATWK
22068	52436	A	22196	2271	3074	RDPSTPTVVRSLGVPLLSRRRFK
22008	32430	^	22190	2271	3074	MAPSVAATKLAERNETMVQTC
						WTETQSSSDGSTPRRVRISLAL
1					}	WRPSLFFSMADTSLVLNFPRVS
1						SSGVIRPSNKRPSLAILTRISRTS
						SPKYMPLTIFSNSCFPGLRDPRSI
						SLSNSVRMSSNDPSSPKAPHS/G
						IDTHPPLVSFHQINVPQFLHVAS
						IGASADDQADFTVLVVIATGHH
						GAHCVIHHSYDVGIIVPSLFDGF
						VKQFDHICSFHSARFKSLCPGD
L	<u> </u>	<u> </u>	<u> </u>		<u> </u>	QDGLIYFGLLARKRKRESHR

SEQ ID	SEQ ID NO: Met SEQ ID NO: Nucleotide		Nucleotide location of last Amino acid sequence (X=Unknown,			
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22069	52437	Ā	22197	1956	2747	FAFSPK*HSCLRCPCI*FSSGLLH
						EVL*LLPLCWP*THGWDPGSRE
						ANKSPKLHAIRCVWVLEEENL
						WLSSNSQSLQTVKN*ESHIN*SC
						RSNLI/HH*FWNQVK*K*LLNIS
						GNCFFFLRWSL/DSVAQAGVY
						WRDLGSLQPPPPGFKRFSCLSLP
		1				SSWDYWHLPPCLAKFCIFSRDG
						/GFTIWARLVLNS*SCDLPASGS
						QSAGITGVSHHTW\LQVTYFLK
						EMRSCYFSQVGWPQTPGLKQC
			ŀ			SHLKLLSSWDYRHMSPHLAISG
22070	52438	С	22198	85	228	
22071	52439	Α	22199	2	434	
22072	52440	Α	22200	1	1356	
22073	52441	Α	22201	146	1909	ILHSLEISALLSLMMLTIGDVIK
	Ì					QLIEAHEQGKDIDLNKVKTKTA
						AKYGLSAQPRLVDIIA\SVPPQY
1						RKVLMPKLKAEPIK\TA\SGIVV
1		ļ				VAVMCKPHRCPHISFTGNICVY
		1				CP/GVGPDSDFEYSTQSYTG\YE
						PTSMRS/IFRARYDPFLQTKDRI
ļ		ļ				EQ*RQLGHSVDKVEFIVMGGTF
	ļ	İ				MALPEEYRDYFIRNLHDALSGH
						TSNNIYEAVKYSERSLTKCIGITI
						ETRPDYCMKRHLSDMLTYGCT
	ļ					RLEIGVQSVYEDVARDTNRGH
	ľ					TVKAVCESFHLAKDSGFKVVA
	ĺ	l				HMMPDLPNVGLERDIEQ\FTEFF
	I					ENPAF\RPDGLK\IYPTLVIRGTG
						LYELWKS\GRYKSYFS**PGLN
		l				WVARFLALVPPWTRVYRVQRD
		l				IPMPLVSSGVEHGNLRELALAR
:		1				NEDLRIQCRDVRTREVGIQEIH
	1					HKVRPYQVELVRRDYVANGG
		1				WETFLSYEDPDQDIL\IGLLR\LR
		1				KCSEETFRFELGGGVSIVRELH
	1					VYGSVVPVSSRDPTKFQHQGFG
						MLL\MEEAERIS*RRNMGLGKI
						AVISGVGTR\NYY\GKIGYRLPR
						PRTMVKMLEIMGHTQSHSFCSI
		L				LPWHEHRRIRISLNTQQRGL
22074	52442	A	22202	1	457	
22075	52443	Α	22203	2	297	CVMYRDYPLELFMAQCY/GNIS
		1				DLGKGRQMPVHYG/CKERHFV
		1				TISSPLATQ/IPQAARGPGYGIM/
		}				SIRVDGNDVFAVYNATK/EARR
		1				RAVAENQPFLIEARPT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	r i	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22076	52444	A	22204	3	1446	DWLLAKMAEAIAAAR\VWRLN RGLSHAALLLLRQLGGSGD*AR SHPPRQQQQCSALDDKP\QFPG ASAGVIDKLEFIQ\PNVISGIPIYR VM\DPQGQIHQPQARDPHLAEG ERWLKL\YKS\MTLL*HPWDRIL **VFSGQGRILLLT*TNYWLRR GTHVG\SARRLGQTRTLVFGQY RE\AGVLMYRDYPLELFMAQC YGNISDLGKGRQMPVNYG\CKE RHFVTISSPL\ATQDPSGRWGAA YAATGGANANRVR/VICYFRAR GAAS*GGDAHA\GF\NFRCHNL SCPIIFFCRNNGYAISTPTSEQYR GDGIAARGPG\YGIMSIRVDGN DVFAVYNATKEARRAVAENQ PFSHQAHDSRVQAPSP\SDAS*T YRSVDEVN\YWENKGPPILPGF GHYLFSQ\GWWDEEQQKAWR KQSRRKDPGPVQPS\VMEAFEQ AERK\PKPNP\NLL\FSDVYQNM PAPVRQPPES\LARHFKTYGNH YPWDHFDK
22077	52445	A	22205	2	292	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22078	52446	A	22206	1	1894	QSLADLLQSQPSITLTSQIWPSP
22070	32710	'	22200		1.05.	A*GFWSWGNLSVTTPSK\PATP
						TRTIAATPIQTLPQSQTAPKRID
						TPSLEEPSDLEELEQFAKTFKOR
						RIILGFT\QGDAGLAMVKLYGN
						DESPTTIFREALNLSFKNMCKL
ŀ						KPLLEKWLNDAENLSSDSSLSS
						PSALNSPGIEGLSRRRKKRTSIE
į						TNIRVALEKSFLENQKPTSEEIT
						MIADQLNMEKEVIRVWFCNRR
						QKEKRINPPSSGGTSSSPIKAIFP
ŀ						SPTSLVATTPSLVTSSAATTLTV
		ļ				SPVLPLTSAAVTNLSVTASTSEA
	ļ					SSASETSTTQTTSTPLSSPLGTSQ
			•			VMVTASGLQTAAAAALQGAA
						QLPANASLAAMAAAAGLNPSL
						MAPSQFAAGGALLSLNPGTLSG
						ALSPALMSNSTLATIQALASGG
						SLPITSLDATGNLVFANAGGAP
						NIVTAPLFLNPQNLSLLTSNPPG
						GFTIEISNNNSTMVMTGMRIQIG
						TQAIERAPSYIEIFGRTMQLNLS
						RSRWFDFPFTREEALQADKKLN
		ŀ				LFTVRKFQNIRHTLFKCENKNT
1						FQQCLLLEQDSSCGTGFCSMTT
						YPVMLLSSEQPVQSLSVRQCVI
						LLDAAAVISGGMGAVCLLLTH
						LCPLCYAGLLSEITDPESFVKDL
						LYSVP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22079	52447	A	22207	1	2417	MRVRTSSLPSTMLRVQNYARA
						KAGPEYAFVDSTECSPQWMVS
						PLKRGTQTNGLDFQKQPVPVG
						GAISTAQAQAFLGHLHQVQLA
						GTSLQAAAQSLNVQSKSNEESG
		1				DSQQPSQPSQQPSVQAAIPQTQ
						LMLAGGQITGLTLTPAQQQLLL
						QQAQAQAQLLAAAVQQHSAS
						QQHSAAGATISASAATPMTQIP
						LSQPIQIAQDLQQLQQLQQNL
						NLQQFVLVHPTTNLQPA\QFIIS
						QTPQGQQGLLQA\QNLLTQLPR
						QSQANLLQSQPRI\TLTSQPATP
						TCTIAATPIQTLPQSQSTPKRIDT
		ŀ		İ		PSLEEP\SDLEELEQFAKTFKQR
				1		RIKLGFTQGDVGLAMGKLYGN
				1		DFSQTTISRFEALNLSFKNMCK
						LKPLLEKWLNDAENLSSDSSLS
						SPSALNSPGIEGLSRRRKKRTSI
l						EA\NIRVALEKSFLEN\QKPTSEE
						ITMIADQLNMEKGVIRVWFCN
						RRQKEKRINPPSSGG\TSSSPIKA
	1					IFPSPTSLVATTPSPVTSSATTTL
						TVSPVLPLTSAAVTNLSVTGTS
	Į					DTTSNNTATVISTAPPASSAVTS
						PSLSPSPSASASTSEASSASE\TS
ŀ						TTQ\TTSTPLSSPLGTSQVMVTA
			}			SGLQTA/AQLLPFKGAAQLPAN\
ŀ	1					ASLA\AMAAAAGLNPSLMAPS
						QFAAGGALLSLNPGTLSGALSP
						ALMSK\STLA\TIQTLASGGSLS
		1				K*TSLECKLGTLVICPMREEPPD
22080	52448	A	22208	3	114	
22081	52449	A	22209	1	505	NLGVDLLPGYQDPYSGRTLTK
						GEVGCFLSHYSIWEEVVARGLA
	Ì					RVLVFEDDVRFESNFRGRLERL
			}			MEDVEAEKLSW\YSYWTLAYA
						LRLAGARKLLASQPLRRMLPV
						DEFLPIMFDQHPNEQYKAHFWP
						RDLVAFSAQPLLAAPTHYAGD
						AEWLSDTETSSPWDDDSGR
22082	52450	A	22210	3	626	RAARAAPLLHLLLLGPWLEA
	152.50			ľ		AGVAESPLPAEVLAILARNAEH
				1	1	SLPHYLGALERLDYPRARMAL
				1		WCATDHNVDNTTEMLQEWLA
						AEGDDYAAVVWRPEGEPRFYP
					1	DEEGPKHWTKERHQFLMELKQ
		1				EALTFARNWGADYI/LGATLGL
						WQDHSVNSLGGCSPGRAGACD
				1		VPSSKFADTDNILTNNQTLRLL
ĺ						MGQGLPVVAPMLDSQTNV
Ļ	<u>L</u>	<u> </u>	<u> </u>	<u> </u>	1	MOGOLI A ANTMINDOGLIMA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22083	52451	Α	22211	1	2004	MRAARAAPLLQLLLLLGPWLE
						AAGVAESPLPAVVLAILARNAE
						HSLPHYLGALERLDYPRARMA
						LCLQGPAQPQWDLWLAGFEEE
						QTGTSEVHWGPTWQGSTTLEA
ĺ						SGHRCATDHNVDNTTEMLQE
						WLAAVGDDYAAAVWRPEGEP
						RFYPDEEGPKHWTKERHQFLM
						ELKQEALTFARNWGADYILFA
						DTDNILTNNQTLRLLMGQGLPE
						VAPMLDSQTYYSNFWCGITPQ
						GYYRRTAEYFPTKNRQRRGCL
ļ						RVPMVHSTFLASLR\AEGADQL
						AFYPPHPNYTWPFDDIIVFAYA
						CQAAGVSVHVCNEHRYGYMN
						VPVKSHQG\LEDERVNFIH\LILE
						ALVDGPRMQASAHVTRPS\KRP
						SKIGF\DEVFVISLG\RRPDRRER
1						MLASLWEMEISGRVVDAVDG
						WMLQQQVPFRNLRP*DLLPGL/
1						YKDPYFGPHS*PRAEVGLLSSD
					:	HYSIW\EEVVSQGAWPR\VLVV
						*RNDVAPLRTNFQGGGLERL\M
						E\DVEGKRNLSWEKPPFRILPLG
}						R\KQVEPLRRET\AVEGLPG\LV
						VAG*LPNW\TLA\YALRLAGGP
	•					ASLLALQPFAPHCCPVERISLPI
ŀ						HCFDQ\HPNEASTKAHFWPRD\
						LVAFSAQ\PLLAAPTHYAG\DAE
						WILLSRRHHSPWDDDSGRLISW
						SGSQKTLRSPRLDLTGSSGHSL
22084	52452	Α	22212	1	468	ELSHNAKAKVMVASHNEDT/V
						LFALRRMEELGLHPADHQVYF
						GQLLGMCDQI\SFPLGQAGYPIL
						TQIHLFSPHTCRAAGGEVRCLP
						ALPRVWALRCGPNLIPAWDSH
						WKLLGTLLELCAQGPHLCDPH
						VLGPRGLSTFCQGQPTQPEPLG
						EQWPGWG
22085	52453	Α	22213	1	1893	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22086	52454	A	22214	390	2005	INRDVETACTYILRNTEDVMLE
						FVMREWKKSRKLLGQRLFNKL
						MKMTFYGHFVAGEDQESIQPL
			•			LRHYRAFGVSAILDYGVEEDLS
						PEEAEHKEMESCSSAAERDGSG
						TNKRDKQYQAHRAFGDRRNG
						VISARTYFYANEAKCDSHMETF
		İ				LRCIEASGRVSDDGFIAIKLTAL
						GRPQFLLQFSEVLAKWRCFFHQ
			;			MAVEQGQAGLAAMDTKLEVA
						VLQESVAKLGIASRAEIEDWFT
						AETLGVSGTMDLLDWSSLIDSR
						TKLSKHF/V*VPNAQTGQLEPLL
			1			SRFTEEEELQMTRMLQRMDVL
						AKKATEMGVRLMVDAEQTYF
						QPAISRLTLEMQRKFNVEKPLIF
						NTYQCYLKDAHDNV\TLD\VDL
						SPREGWCFGAKPVRGAYLAQE
		ĺ				RACAAEIGYEDPINPMYEATNT
		l				MYHRCLDYVLEELKHNAKAK
						VMVASHNEDTVRFALRRMEEL
						GLHPADHQVYFGQLLGMCDQI
						SFPLG\QAGYPVYKYVPYGPVM
						EVLPYLS\RRALE\NSSLMKGTH
			:			\RER\QLLWLELLRRLRTGNLFH
22087	52455	A	22215	3	303	WELLOW EDEDLICE GIVEN IN
22088	52456	A	22216	798	2250	VPRAPQGML*GPRKDAGTPER
						RAGPPGGRGTADGGGCGALAG
						SGS/DDPRLCIWTFLVTDQLLLP
						TGPGRPRGSVAAAATGPGVGA
						LRAPARAQPRGAVDGRPLLPQ
		ĺ	}			VLPHRRLEVSLGKSPKECVLW/
						RHLLSESEKRPFVEEAERLRVQ
						HKKDHPDYKYQPRRRKSAKAG
						HSDSDSGAELGPHPGGGAVYK
						AEAGLGDGHHHGDHTGQTHGP
						PTPLTTPKTELQQAGAKPELKL
						EGRRPVDSGRQNIDFSNVDISEL
						SSEVMGTMDAFDVHEFDQYLP
				,		LGGPAPPEPGQAYGGAYFHAG
						ASPVWAHKSAPSASASPTETGP
						PRPHIKTEQPSPGHYGDQPRGSP
						DYGSCSGQSSATPAAPAGPFAG
						SQGDYGDLQASSYYGAYPGYA
						PGLYQYPCFHSPRRPYASPLLN
						GLALPPAHSPTSHWDQPRRNGS
						TAQGPKEGVAGALQGAVRSKA
		1				GLGAEKPLHSRPHSLDNGCTGC
l	ì	ı	l	1_	l	PGSGVSVRRLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
]	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \≔possible nucleotide insertion)
]				sequence		
22089	52457	IA	22217	299	1631	LLEPHCGGKCFKRHWFPCSCEA
22007	32437	`	22217	277	1031	SC\PSAAPQRW*HTG/SPEVPAA
		ŀ				PGAQGREPSM/RDGEGAVRGGS
1						CMEAGPMGLQCVDKKHQGVH
	:	ŀ				N/RGEARVFLQSQVGAILPRPW
į						PNPAPSEP/VQSPLASSTGP*VFP
1						SRPGQQASGPA/RRHPGAEPGQ
						RGDAVQLLPRPQTSPHGPQDA
	1					ALSSAPHPWRQL/MASPGLQEA
						PLHPFCGHRSPCPVPWQLSPTD
						PIQSRPPARGWAPANQGRPSHP
						RCRGGGGAFCGHGF/SYTOPLT
]				·		FRLASMVLSIHPKPPKETLRFRP
						NMVVAAEPGVPAMSQRRPRGC
1						TRFSHTDKAPLDTCTDMNAYT
į.						CARRHSSTYSDHITLQGAFACT
						1
i						HTWAQSCTCQPPPTPAAKCLDR
						GQSSGMLHAGCVGWGGGTHV RVDLSSQLNRRVRPHLCMLRC
						l '
						WSSCHRASVSCTHPSLHTTGPV LISARTGADHTWYTAGTY
22090	52458	A	22218	383	762	RHQGNQKRKELSLK/CQKSRQE
22090	32438	A	22218	363	702	SPRAEENPKWREGKKETSESSV
		l				QKAGRAAAAQAGAAASRVPG
						LSGSNLAPCNKGRLSAREDVSN
		ŀ				SKLKILSSGRKHTEHVIMLKAW
22201	50450	ļ	22210	205	0.45	GLGRQSLRCRLFDLIPSHKV
22091	52459	Α	22219	325	945	QPSPLAPKADLSAREGRFQLKV
						RCRLQQYQQR\RKF\AAAFLA
		ŀ				FHFSLLGSCGILAEAG\KKEKPE
						KK\VKKSDCGRIGSWSVCVAPP
						SGDCGLGHHGRATRT*\AECKQ
						\TMKTQRCKIPCNWEGRQFGRG
						SAKLPSFQAWGECDLDT\ALKT\
						KTGSLKRALHKSPNAQKTVTIS/
						KSPCGKLTKPQNLQAES\KKKK
	1					KEGKKPEGRLRD

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22092	52460	A	22220	1	1908	MKYIQDFQREKQEFERNLARFR EDHPDLIQNAKKSDIPEKPKTPQ
	1					QLWYTHEKKVYLKVRPDATTK
	1					EVKDSLGKQWSQLSDKKRLK
		-				WIHKALEQRKEYEEIMRDYIQK
	[ļ				HPELNISEEGITKSTLTKAERQL
		j				KDKFDGRPTKPPPNSYSLYCAE
	ļ	İ				LMANMKDVPSTERMVLCSQQ
						WKLLSQKEKDAYHKKCDQKK
	İ	1				KDYEVELLRFLESLPEEEQQRV
•		İ				LGEEKMLNINKKQATSPASKKP
		ļ				AQEGGKGGSEKPKRPVSAMFIF
		ļ				SEEKRRQLQEERPELSESELTRL
						LARMWNDLSEKKKAKYKARE
						AALKAQSERKPGGEREERGKLP
						ESPKRAEEIWQQSVIGDYLARF
						KNDRVKALKAMEMTWNNME
						KKEKLMWIKKAAEDQKRYERE
		ļ				LSEMRAPPAATNSSKKMKFQG
	1					EPKKPPMNGYQKFSQELLSNGE
						LNHLPLKERMVEIGSRWQRISQ
		•				SQKEHYKKLAEEQQKQYKVHL
		ļ				DLWVKSLSPQDRAAYKEYISN
	1					KRKSMTKLRGPNPKSSRTTLQS
					1	KSESEEDDEEDEDEDEEEE
	1					DDENGDSSEDGGDSSESSSEDE
	١ ،					SEDGD/EGGADRAQWEGRGVL
						PGGPPGDSTSFLTPQNEEDDED
						EDDDEDDDEDEDNESEGSSSSS
		ļ				SSSGDSSDSDSN
22093	52461	Α	22221	129	2365	DOL DELLOCE WIEL A LEAGUE
22094	52462	A	22222	5	2730	PSLRFAISGFVWRLAAPASAAA
						ELRLDCSARRAPRCPRPALPSTR
						SAEGAVLCAFRREEQQAAATA
		•				AAAAAATAAAAAAPAPPPRP EEEPLPPREGAAAVPGRAGEGA
						ATQSQGGSRWSGKPGGRAAPA
	İ	ł				RPRRWLDSWRMNGEADCPTDL
		l				EMAAPKGQDRWSQEDMLTLLE
						CMKNNLPSNDSSKFKTTESHM
						DWEKVAFKDFSGDMCKLKWV
						EISNEVRKFRTLTELILDAQ*HF
	1					KNPYKG\KKLKKHPDFPK
22095	52463	Α	22223	1	446	RPSPPTSSM/PGLWGYSRPCSSG
[1	`		1	1.10	VPASALQA*SPAFSPASQRPPTT
						SWWEVP*EEGRA\PVWEGRIRA
					1	VGPAPAS*PSDLPAWSCRSWRE
					1	TRG\SPAEDAMLLRAVEPLEGC
						QRPLLDSEPP/SPARTSAPAASA
						`
						TRG\SPAEDAMLLRAVE

SEQ ID	SEQ ID NO:	1	I .	l l		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22000	I52464	<u> </u>	122224	1582	1175	 THTHVPAGTHPHTQTTSHSHLI
22096	52464	A	22224	382	11/3	GRALTVPESPKCPCLLPSAC*PR
						PGRAQPAGPAWSKKEASNPGS
						E/PPQHSL*RRMAPPE\PVPARA
						VHTCRCLSSAPPGARLPLDPER
:		İ				RAPQSPRSPPPSAQPPPPPPPPPP
						PPPGRRGPSQQSWRPSPPPPRTR
		1				SRSPRRSRVRGPRAPALAAPPPP
İ						GARPGPCAAACGSPTGCS
22097	52465	A	22225	609	5241	GPQTTWCGRPTMCQHCGPAVS
220)	32403	``				SWKQPSHDPCCRGGENRRQPG
ŀ		1				TPARATQGRLRRRPEGARVPSA
	-					RRGRPRARWRDKERNNTCVA
İ						ML*QVTNSTAN\APQNNPEVDS
ŀ					İ	IMAELKVEAPASVDWQKRCLT
ļ						LETQLFRFRLQASKIRELLADK
						GHGTYISPFSLAKQMQELEQRL
						LEAEQRAENAETQVGVMEEKV
						KLSNLKNVDSEGSLHRKYQELL
						KAIKGKDELISQLEAQLEKQKQ
						MRAEEAKTVQEKAAKIKEW
22098	52466	A	22226	2	205	
22099	52467	Α	22227	1	6186	
22100	52468	Α	22228	96	1584	DLGSGITASPPRPEPEHPVFPPA
						PHTPSKMGKEKTHINIVV\IGHV
]	\DSGKSTTTGHLIYKCGGIDKRT
						IEKFEKEAAEMGKGSFKYAWV
						LDKLKAEREPG\ITIDISLW\KFE\
						TTKYYITIID\APGHRDFIKNMIT
		1				G\TSQADCAVLIVAAGVGEFEA
		1	:			GISKNGQTREHALLAYTLGVK
		1				QLIVGVNKMDSTEPAYSEKRY
1		1				DEIVKEVSAYIKKIGYNPATVPF
		1				VPISGWHGANMLEPSPNMPWF
		1				KGWKVERKEGNASGVSLLEA\
						LDTILPPTRPTDKPLRLPLQD\V
ļ		1				YKI\GGIGTVPVGRVETGILRPG
		1				MVVTFAPVNITTEVKSVEMHH
						EALSEALPGDNVGFNVKNVSV
		1				KDIRRGNVCGDSKSDPPQEAAQ
						FTSQVIILNHPGQISAGYSPVIDC
						HTAHIACKFAELKEKIDRRSTK
		İ				KLEDNPKSLKSGDAAIVEMVPG
		1				KPMCVESFS\Q*PVIG/RTPVRD
						MRQTAA\VGVIKNVQKKSVGA
						GKVTKSAQKAQKAGK
22101	52469	Α	22229	1	1023	
22102	52470	Α	22230	1	396	
22103	52471	Α	22231	2	1259	
22104	52472	Α	22232	1	2514	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22105	52473	Α	22233	3	2027	EESDCLTEYEEDAGPDCSRDEG
		l				GSPEGASPSTASEMDVSLHFVL
		1				WGCLHVYQRMIDKAEDVCLFV
						AQPGELVGQLAVLTGEPLIFTL
		1				RAQRDCTFLRISKSDFYEYDSP
		}				KLECGWHSEDPNPWTSPSTGL
		ļ				NPLFLELCRHMVFQRLGQGDY
		1				VFRPGQPDASIYVVQDGLLELC
		ł				LPGPDGKECVVKEVVPGDSVN
						SLLSILDVITGHQHPQRTVSARA
		ŀ				ARDSTVLRLPVEAFSAVFTKSA
						RTITGCVWRYVRASASYCPYLP
						PLCDPKDGHLLVDGCYVNNVP
						GSLWRYVRASMT\RRATCPRCA
						TPRTGTYSWMAATSTICQASGR
						PHHPHTQAPPAPHTRTRRTSPA
						AWVPKRSSPLTWGARMRRTSA
						PTGTACPAGGCCGSG*IPGLTR*
					1	RFQTWLKSSPAWPTCPVCGS*R
				·		LSSPAPTASTCARPSTASRPWTL
			ĺ			GSSTRSILRTTRRMSTAVTTTAT
,						PAPIITPST*GTNGTGCAPQIRPA
						EPSPGARTSWNPSLSATFAPEEF
į						VASP*LDVPICRMGQCCSLPGG
		1				ATLP*GPCGPQCMPGNVVPPER
						TAARYRQTTAYQTPACPTPPRG
						QRPT/CVPTVSSLQAGLQGGPRS
						DFDMAYERGRISVSLQEEASGG
						SLAAPAREPREQPAGACEYSYC
						EDESATGGCPFGPYQGRQTSSIF
			1			EAAKQELAKLMRIE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22106	52474	A	22234	281	1832	NDSGLPDRPSSWNQPMEAPLQ
22100	32474	,	22254	[20.	1032	TGMVLGVMIGAGVAVVVTAV
						LILLVVRRLRVPKTPAPDGPRY
						RFRKRDKVLFYGRKIMRKVSQS
						TSSLVDTSVSATSRPRMRKKLK
						MLNIAKKILRIQKETPTLQRKEP
						PPAVLEADLTEGDLANSHLPSE
				:		VLYMLKNVRVLG\HFEKPLFLE
						LCRHMVFQRLG/QGDYVFRPG
						QPDASIYVVQDGLLELCLPGPD
						GKECVVKEVVPGDSVNSLLSIL
						DVITGHQHPQRTVSARAARDST
						VLRLPVEAFSAVFTKYPESLVR
						VVQPGPPSGPAPYPAPSYGHAP
						RALQIIMVRLQRVTFLALHNYL
						GLTNELFSHEIQPLRLFPSPGLPT
						RTSPVRGSKRMVSTSATDEPRE
						TPGRPPDPTGAPLPGPTGLQGG
						PRSDFDMAYERGRISVSLQEEA
						SGGSLAAPAREPREQPAGACEY
						SYCEDESATGGCPFGPYQGRQT
						SSIFEAAKQELAKLMRIEDPSLL
						NSRVLLHHAKAGTIIARQGDQ
						MEEVKVPAFAVIVD
22107	52475	A	22235	50	4287	PGRKSPGAEDRALWGPACRLL
						TDSGLPDRPSSWNQPMEAPLQT
						GMVLGVMIGAGVAVVVTAVLI
						LLVVRRLRVPKTPAPDGPRYRF
			ł			RKRDKVLFYGRKIMRKVSQSTS
						SLVDTSVSATSRPRMRKKLKM
						LNIAKKILRIQKETPTLQRKEPP
						PAVLEADLTEGDLANSHLPSEV
			i S			LYMLKNVRVLGHFEKPLFLELC
						RHMVFQRLGQGDYVFRPGQPD
						A\SIYVVQDGLLELCLPGPDGKE
						CVVKEVVPGD\SVNS
22108	52476	A	22236	283	403	TCLSLVSGCGHCEKMNSEENW
						*IPIQWASSNPGRRRKNIR
22109	52477	Α	22237	33	222	NGPM\CFVKQLEIPQYGYRNNV
						PTTTPRSNLAKELEKYSKTSFEY
L						TINDNHTYGGLGCMRPRLV
22110	52478	В	22238	1	1860	
22111	52479	A	22239	303	672	GITACVFFSMAEGRTRAERSCD
				}		SK*AIGAPAEE*GGASPGKSPEL
						WQGAGYSPLGACGHKARLWIP
						WRRSWSGEKGARGCRGTGKK
						LGHQGWRWETPGRPGGAGPA
			ļ			GFPDPVVMGCEKEASRV
22112	52480	Α	22240	1	1377	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22113	52481	Α	22241	1	2232	MQDTEKDDNNNDEYDNYDEL
						VAKSLLNLGKIAEDAAYRARTE
						SEMNSNTSNSLEDDSDKNENLG
						RKSELSLDLDSDVVRETVDSLK
		ŀ				LLAQGHGVVLSENMNDRNYA
						DSMSQQDSRNMNYVMLGKPM
						NNGLMEKMVEESDEEVCLSSL
						ECLRNQCFDLARKLSETNPQER
						NPQQNMNIRQHVRPEEDFPGRT
						PDRNYSDMLNLMRLEEQLSPRS
						RVFASCAKEDGCHERDDDTTS
		ŀ		1		VNSDRSEEVFDMTKGNLTLLE
						KAIALETERAKAMREKMAMEA
						GRRDNMRSYEDQSPRQLPGED
						RKPKSSDSHVKKPYYGKDPSRT
						EKKESKCPTPGCDGTGHVTGL
						YPHHRSLSGCPHKDRVPPEILA
						MHESVLKCPTPGCTGRGHVNS
						NRNSHRRDIQVFNLQKQEMIDR
						LESLSGCPIAAAEKLAKAQEKH
						QSCDVSKSSQASDRVLRPMCFV
						KQLEIPQYGYRNNVPTTTPRSN
						LAKELEKYSKTSFEYNSYDNHT
						YGKRAIAPKVQTRDISPKGYDD
						ANGYCKDPSPSSSSTSSYAPSSS
						SNLSCGGGSSASSTCSKSSFDYT
						HDMEAAHMAATAILNLSTRCR
						EMPQNLSTKPQDLCATRNPDM
						EVDENGTLDLSMNKQRPRDSC
						CPILTPLEPMSPQQQAVMNNRC
						FQLGEGDCWDLPVDYTKMKPR
						RIDEDESKDITPEDLDPFQEALE
22114	52482	Α	22242	727	4372	LWACVLPGVTLPPFLGEKTSCK
						MEVDTEEKRHRTRSKGVRVPV
						EPAIQELFSCPTPGCDGSGHVSG
						KYARHRSVYGCPLAKKRKTQD
						KQPQEPAPKRKPFAVKADSSSV
						DECDDSDGTEDMDEKEEDEGE
						EYSEDNDEPGDEDEEGDRE
		Ì				EEEEIEEEDEDDDEDGEDVEDE
						EEEEEEEEEEEEENEDHQMN
						CHNTRIMQDTD\LDDNNNDEY
			:			DNYDELVAKSLLNLGKIAEDA
		<u></u>				AYRARTESEMNSNTSNS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
}				sequence		
22115	52483	A	22243	34	1558	PASEEFNSLHGRTPRRDLECRN
ł						QQSTTEERRRHDCVQQASEHN
1						EGKWDFAFSAQSQPKP*KPFDV
						PC*DSRPACP*EHTAGTSGSTSS
						ETCFFQSSEWPYGELGAYNIFT
						GPKSTLSSLVTGVSNSSSLS*LP
						SPKNTITNWFHTRDHSSYSEIHS
						KACANP*APSQPSR*ASW\IATG
					•	SFRRAGLAT*PCCTGSKLLPAW
						VLANHRWDRTRDGIQKQGNSE
						*PSSPGTRAFEGNSSSPAPAASIT
						SMLSPSFTPTSVIRKMYESKEKS
						KEEPASGKAALGDSKEDTQKA
						SEENLLSSSSVPSADRDYFSHY
						KFQTVRHYRGSSCS\PHLSQGQP
						FTPKEQDYSDLKGNWGRKNTH
						FGHSPVSLQHFFFRPVPPSSPLSP
						MVPNG*GLLTKLHPGFGTEDA
						GPGSTSTASSKFAPNWCASSWD
						GLESFTGNIWPHPGSALLPFN*S
						LVHPLLNPRPGNTSASGNGATA
]						ANSAQ\SCILQALVSHGSSCSAF
						QDNPSETCPAGQACPHALP
22116	52484	A	22244	1502	2657	EGKVPLAFIVLTSLLNAVMSPSF
						FSCRLLVTALKVSSRCSAMKGV
						EFFTCWSTFKPFRPASTSSTEST
						TPECELSLSFSLFAERLERRGFK
Ì						STLAFCSISKISTLFSAWSSNGIG
						AK*FPEFCPGDRMACSRPASLSS
						SSCG/S*PKTAGSAS*SARVREPP
						TEPTGRD*PFSQ\TSSIIEAKHGT
						LSRLKNSLKSKSPGDCSGKTAS
						LGTS*SAAGSCARMTSTSSSSAT
						PPLHSTIPSFTEAVRRLVLFLPL*
						SSSSILSSKPVSSMVSD*LVGPA
						ENHSGSSSV*ESFLLRSPKTRLL
						SPNSLLKRLSLKSLSRSLSLKSR
						RSLSL\SGDPSQRSGQRLSSHCQ
						SYVHQADEAHHYLSLMDFLSA
						GS*QRLSRGSPLQSSVSEG
22117	52485	Α	22245	324	391	
22118	52486	Α	22246	42	274	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22119	52487	A	22247		835	MCAGNLCSWEERWAVDEKEA TDKKPHVGGFGQGAPAPAAPA AAMSGRSVRAVTRSRF*DDIKK VMAAIEKVRKWEKKWVTVGD TSLRIFKWVPVTDSKEKEKSKS NSSAAREPNGFPSDASANSSLL LEFQDENS\NQS\SVSDVYQLKV DSSTNSSPKPPKQSESLSPAHTS DFRTD\DSQPPTLGQGDPGRSPP LP\SSEVADEPPTLTKGRTSFHL ETQVVEE\EEDSGAPPLKRFCV\ DQPTVPQTGSEKLAPSRPSAFW ALPLFIAFWFWPRRVCWG
22120	52488	Α	22248	3	244	
22121	52489	В	22249	271	711	
22122	52490	В	22250	73	200	
22123	52491	A	22251	885	975	GSTWWPAD*AQL*LHLPTRCM AECWEIPA
22124	52492	Α	22252	102	909	PGGRHGRQNDFAAPGGRLEARI HGTVSAPLYHRVLCRLLGSRPA SAAGPAARGGDLAPGEHSFCE DGNYARVESVDRSAINRAMQN PRKSTASVCWTFTASDLPAQGV \PRTSSTRTLREDPFPPSAEKWL RAVLHHFVNEKLQQIFIELTLKP SRRSMSGRHPLDSNPVLQQQGR CDLIENKLSPPGIMSVLDDVCPP CTPRAGEQTRHCCRTAGGCGD PRAFQQLERRFVIHHYAGKTHR PLPGLTPTGNSTNARNIVMDTQ QRTGNN
22125 22126	52493 52494	A	22253 22254	48 1243	112 1615	YSPSSWFSNLYQPSTGALSSSET PKDMAPMKSLREVRSQSHTSTL SRRSTSPGFRLNTKVSFHTGFSV AFTVRVFCFVRFRGSGMSFSFT* GSERPLGSIGIRLRASPTVTWISS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22127	52495	A	22255	468	11431	SVSTTRSFSVDSSAKTAAMPVT
22.2.	32.55			1.00		VTRTTHHNHPDVIFGAWGPP*S
						WGPLWALTQPLGLL\RLLQL\V
						STCVAFSLVASVGAWTGSMGN
						WSMFTWCFCFSVTLIILIVELCG
						APGPFPPWSWR\NFPI\TFACYA
						ALFCLSASIIYPTTYVQFLSHGR\
						SRDHAIAATFFSCIACGC\YATE
						VAWTRARPGEITGYMATVPGL
						LKVLETFVACIIFAFISDPNLYQ
						H\QPALEWCVAVYAICFILAAIA
						ILLNLGECTNVLPIPFPSFLSALA
						\FCLSSSMPPPLFSGPSTSSMRSM
		1				AASLGAREM*AAAAGMPTTW
						VPGTADWLRPS
22128	52496	A	22256	98	414	CTRVSVAEKTSVCLFGLYWESL
22120	32490	^	22230	36	414	SISSSCESSAKWPFSTIRSASSTT
					}	RHLSCRRFLK**CFSLINSHSRP
						GVATTISGRRFSILCCFCVDIPPT
						KIAVLIPVNLAT
22129	52497	A	22257	2	224	RIAVEIFVNEAT
22130	52498	A	22258	2	115	
22131	52499	A	22259	2	118	
22132	52500	A	22260	472	648	GPGGRSFTVWWNW*RRMAQC
						P\PAEGVQRPECLGPHRAVPAP
						AHVGLHQEPSRLLPALH
22133	52501	Α	22261	540	1893	GRCATPRPRWRGHQCDPPPGD
						VCVAWREMQLRNKKRRR\DLG
						NLIK\RFLTAPNSLPD/AVVLTTE
						EAVIFWCNVLAQQIFGL/PLEM
						MNEQPLEGAVREKALHTMREQ
						TQRMEGLVKQL/LTLSKIEAAPT
						HLLNEKVDVPM/MLRVVEREA
						QTLVYNAVNHTPEGTHITVRW
						QRVPHGAEFSVEE*SEGWPVPV
						PKPHRAVVGANAHISFCPPQEL
						SKRFTAIRKTKGDGNCFYRALG
						YSYLESLLPDAFQPLLPPQAWV
		1				SPGESQGAGIRDPLWPVPVPKP
						HRAVVGANAHISFCPPQELSKR
						FTAIRKTKGDGNCFYRALGYSY
		1				LESLLPDAFQPLLPPQAWVSPG
						ESQGAGTRDPLWLWLFRLWLL
					1	RLLCLLLLLGETAQSRLALTW
						GRHKQGRLARLSGILGFLALRK
						QWLGQQVAQLLLQNLHLLLLR
					}	WAGLGVIVFQVSNQPAWERTG
22134	52502	Α	22262	1	1772	
22135	52503	Α	22263	107	294	LHLLSFGYLTHQLLCLHLKWDS
]			HTLHHSHHTIAR*AGCSEATWS
]	PPRRLRAPRSRALCSGFR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22136	52504	Α	22264	1	474	
22137	52505	A	22265	1	148	VKYHEFTCSFKKQITVPEDARL LK*DVRGRPSRLFLADPPPPHT VNREY
22138	52506	A	22266	509	2755	YQSQSFDEDEMNQVTDWVDPS FDDFLECSGVSTIT\ATSLGVNN SSHRRKNGPSTL\ESSRFPARK\R GNLSSLEQIYGLENSKEYLSENE PWVDKYKP\ETQHELAVHKKKI E\EVETWLKAQVLGRQPKQGG SILLITGPPGCGKTTTLKILSKEH
						GIQVQEWINPVLPDFQKDDFKG MFNTESSFHMFPYQSQIAVFKE FLLRATKYNKLQMLGDDLRTD KKIILVEDLPNQFYRDSHTLHE VLRKYVRIGRCPLIFIISDSLSGD NNQRLLVPKEIQEEWSISNIRFN PWAPTIMMKFLNRIVTIEANKN
						GGKITVPDKTSLELLCQGCSGDI RSAINSLQFSSSKGENNLRPRKK GMSLKSDAVLSKSKRRKKPDR VFENQEVQAIGGKDVSLFLFRA LGKILYCKRASLTELDSPRLPSH LSEYERDTLLVEPEEVVEMSHM PGDLFNLYLHQNYIDFFMEIDDI VRASEFLSFADILSGDWNTRSL LREYSTSIATRGVMHSNKARGY AHCQGGGSSFRPLHKPQWFLIN KKYRENCLAAKALFPDFCLPAL
	·					CRQTQLLPYLALLTIPMRKSSFR FSFYSKDIGEGLPSGRRTLWGRI GKWEGPGLDRGTWDD*TPDSG DEA\QLNG\GHSAEESLGEPTQA TVPETWSLPLSQNSASELPASQP QPFSAQGDMEENIII\EDYESDG DIEASLLIRLLLHSFHFCFHFSGT
22139	52507	Α	22267	2	161	
22140	52508	A	22268		599	VTRLFFQPGYGTLQPQRLLMPK KNR\IAIYE\LLFK\EGVMVPKK\ DVPIPKHPELA\DRNVPNLHVM KPMQ\SLKS\RGYVK\EQFAW\R HFYWYLTK*GYQGVSISSRDYL \HLPP\EIVPCHPYGRSPSRRLAR PSG*KVLEG*ANLRGLHKNGEA DKRLPTRRECLCHPGA\DKKAE AG/ALGSSNPNFQFRGGFGRGR GQ\PPQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22141	52509	A	22269	a t	345	KHPELADKNVPNLHVMKAMQ SLKSRGYVKE\DYLHLPPEIVPA TLRRSRPETG/RPRPKGLEGERP ARLTRGEADRDTYRRSAVPPG ADKKAEAGAGSATEFQFRGGF GRGRGQPPQ
22142	52510	A	22270	2	161	
22143	52511	A	22271	2	641	RRPQHRACSFFFSPGTGPCSRRR CLMPKKNR\IAIYE\LLFKEG\VM VAKKDVHMP\KHPE\LAD\KNG APTFHVMKGHAGLSKSPRATV KGTSFAWGDIFYW\YL\TN*GY QGVSISSRDYLSSCPREICCLAT ATARSPSKRLGKAFGLKRSWE GLRRPGEDFTRGGKFDRGYLT GRIAVATLVADKKAEAG/ALGS SNPNFQFRGGFGRGRQ\PPQ
22144	52512	A	22272		1693	MFNSVPVQNEDGSFNFEGVKA EFRPGTQTQEYIKGMEDSASEV TVNREVTTDNPYTISVTNKTLS AIRIKMFMPR\AYELKVTVIKM AFELSTRCSKRLM/DGSFETVLT DVIEGKTMSGYDRSRRVNLPNF NNQVIFRVVRKTPDSNDSNVV DAIQVRSYAEVIDAKFRYPLTG LLFVEFDSKMFPNQLPTISIRKR WKIVNVPSNYDPESRTYNGNW DGTFKKAWTNNPAWVLYDLM INQRYGLDQKELGISVDKWAL YEAAQYCDQMV\LTAKAARNL ATFVT/CIIQSQTDAPREPAYIFT NDNVVNGDFSYT\SQARRACTR RAT*CLMMSKTCINRTLSQYSI/ V/EGHSTVR\NNVTSITAIG\AHV EA/KANRRGRWILKTNLRSTTV NFATGLEGMIPTIGDVVAISDNF WSSNLTMNLSGRLLEVSGSQIF LPFRVDARAGDFIIVNKPDGKP VKRTISSVSADGKTIEINIGFGFP VKPNTVFAIDRTDLALQQYVVT KIDKGDDDEEFT\SKSRRWSTIL ISTMKLITGLTSTTDRQASLSQI R/CPKPENVK\CPQSRESSRG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22145	52513	Α	22273	1	1843	MEDNLISINKIKILLSVSDGEIDE
						TFSLKQLMFNSVPVQNEDGSFN
						FEGVKAEFRPGTQTQEYIKGME
						DSSSEVTVNREVTTDNPYTISVT
						NKTLSAIRIKMFMPR\AYELKVT
						VIKMAFELSTRCSKRLM/DGSFE
İ						TVLTDVIEGKTMSGYDRSRRV
						NLPNFNNQVIFRVVRKTPDSND
						SNVVDAIQVRSYAEVIDAKFRY
		:				PLTVLLFVEFDSKMFPNQLPTIS
						IRKRWKIVNVPSNYDPESRTYN
						GNWDGTFKKAWTNNPAWVLY
						DLMINQRYGLDQKELGISVDK
						WALYEAAQYCDQMV\LTAKA
ı						ARNLATFVT/CIIQSQTDAYKVV
						RDICSIFRGMSFWNGPREPAYIF
						TNDNVVNGDFSYT\SQARRACT
						RRAT*CLMMSKTCINRTLSQYS
						I/V/EGHSTVR\NNVTSITAIG\AH
						VEA/KANRRGRWILKTNLRSTT
						VNFATGLEGMIPTIGDVVAISD
					,	NFWSSNLTMNLSGRLLEVSGSQ
						IFLPFRVDARAGDFIIVNKPDGK
1						PVKRTISSVSADGKTIEINIGFGF
]						PVKPNTVFAIDRTDLALQQYVV
						TKIDKGDDDEEFT\SKSRRWSTI
						LISTMKLITGLTSTTDRQASLSQ
						IR/CPKPENVK\CPQSRESSRG
22146	52514	В	22274	1	1452	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22147	52515	A	22275	1	1902	MIQKVISGSKGGSQKPHNPVEM
						EDNLISFNKIKILLAVSDGEIDET
1						FSLKQLMFNSVPVQNEDGSFNF
						EGVKAEFRPGTQTQEYIKGME
						DSSSEVTVNREVTTDNPYTISVT
1						NKTLSAIRIKMFMPR\AYELKVT
1						VIKMAFELSTRCSKRLM/DGSFE
1						TVLTDVIEGKTMSGYDRSRRV
						NLPNFNNQVIFRVVRKTPDSND
						SNVVDAIQVRSYAEVIDAKFRY
						PLTGLLFVEFDSKMFPNQLPTIS
						IRKRWKNVNAPSNYDPESRTY
			:			NGNWDGTFKKAWTNNPAWVL
						YDLMINQRYGLDQKELGFSVD
						KWALYEAAQYCDQMV\LTAK
		ŀ				AARNLATFVT/CIIQSQTDAPRE
						PAYIFTNDNVVNGDFSYT\SQA
		l				RRACTRRAT*CLMMSKTCINRT
						LSQYSI/V/EGHSTVR\NNVTSIT
						AIG\AHVEA/KANRRGRWILKT
					ł	NLRSTTVNFATGLEGMIPTIGD
						VVAISDNFWSSNLTMNLSGRLL
						EVSGSQIFLPFRVDARAGDFIIV
		1				NKPDGKPVKRTISSVSADGKTI
						EINIGFGFPVKPNTVFAIDRTDL
		1				ALQQYVVTKIDKGDDDEEFT\S
		l				KSRRWSTILISTMKLITGLTSTT
						DRQASEVKPVNFTSSRHQAQPE
						VGLIPPPLWPQGKGIIAQTSKSS
22148	52516	A	22276	1	675	
22149	52517	A	22277	155	700	SGVLVGHALTARQPSSGRRLV
					:	VDHSANRRGRWILKTNLCSTT
ŀ						VNFATGLEGMIPTIGD/GDFIIVN
ľ						KPDGKPVKRTISSVSADGKTIEI
ł						NIGFGFPVKPNTVFAIDRTDLAL
						QQYVVTKIDKGDDDEEFT\SKS
						RRWSTILISTMKLITGLTSTTDR
						QASLSQIR/CPKPENVK\CPQSRE
			1			SSRG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22150	52518	A	22278		1871	MEDNLISINKIKILLAVSDGEIDE TFSLKQLMFNSVPVQNEDGSFN FEGVKAEFRPGTQTQEYIKGME DSSSEVTVNREVTTDNPYTISVT NKTLSAIRIKMFMPR\AYELKVT VIKMAFELSTRCSKRLM/DGSFE TVLTDVIEGKTMSGYDRSRRV NLPNFNNQVIFRVVRKTPDSND SNVVDAIQVRSYAEVIDAKFRY PLTGLLFVEFDSKMFPNQLPTIS\ IRKRWKIVNVPSNYDPESRTYN GNWDGTFKKAWTNNPAWVLY DLMINQRYGLDQKELGISVDK WALYEAAQYCDQMV\LTAKA ARNLATFVT/CIIQSQTDAYKVV \RDICSIFRGMSFWNGESISVIID RPREPAYIFTNDNVVNGDFSYT\ SQARRACTRRAT*CLMMSKTCI NRTLSQYSI/V/EGHSTVR\NNVT SITAIG\AHVEARPTDCGRWILK TNLRSTTVNFATGLEGMIPTIGD VVAISDNFWSSNLTMNLSGRLQ EVSGSQIFLPFRVDARAGDFNIV NKPDAKPVKRTISSVSADGKTI EINIGFGFPVKPNTVFAIDRTDL ALQQYVVTKIDKGDDDEEFT\S KSRRWSTILISTMKLITGLTSTT
						DRQASLSQIR/CPKPENVK\CPQS RESSRG

NO:				Nucleotide	ruciconac rocarion or mor	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22151	52519	A	22279	3307	5305	CRSSQNTQETSPDRALLVNLRP
						SQGPLLALWVPAEGVGGKGHC
						LPPRSFFRMDNQLVPAVSDGEI
						DETFSLKQLMFNSVPVQNEDGS
						FNFEGVKAEFRPGTQTQEYIKG
						MEDSSSEVTVNREVTTDNPYTI
						SVTNKTLSAIRIKMFMPR\AYEL
						KVTVIKMAFELSTRCSKRLM/D
						GSFETVLTDVIEGKTMSGYDRS
						RRVNLPNFNNQVIFRVVRKTPD
						SNDSNVVDAIQVRSYAEVIDAK
						FRYPLTGLLFVEFDSKMFPNQL
						PTISIRKRWKIVNVPSNYDPESR
						TYNGNWDGTFKKAWTNNPAW
				•		VLYDLMINQRYGLDQKELGISV
						DKWALYEAAQYCDQMV\LTA
						KAARNLATFVT/CIIQSQTDAYK
						VVRDICSIFRGMSFWNGESISVII
						DRPREPAYIFTNDNVVNGDFSY
						T\SQARRACTRRAT*CLMMSKT
İ						CINRTLSQYSI/V/EGHSTVR\NN
						VTSITAIG\AHVEA/KANRRGRW
						ILKTNLRSTTVNFATGLEGMIPT
ŀ						IGDVVAISDNFWSSNLTMNLSG
						RLLEVSGSQIFLPFRVDARAGD
						FIIVNKPDGKPVKRTISSVSADG
						KTIEINIGFGFPVKPNTVFAIDRT
						DLALQQYVVTKIDKGDDDEEF
						T\SKSRRWSTILISTMKLITGLTS
						TTDRQASLSQIR/CPKPENVK\CP
						QSRESSRG
22152	52520	Α	22280	3	412	PAAVA*VKTEPETPGPSCLSQE
						GQTA\VKTEESSELGNYVIKIDH
						LETIQQLLTAVVKKIPLITAK\W
						QRAMTMRKVLQEILEKNPRFH
						HLTPLKTKHIAHWCRCHGYTPP
						DPESLRNDGDSIEDVLTQIDSEP
22153	52521	Α	22281	2	513	IFACWGKQAWPACCNSLRGQA
						VKGNQLLPVSLVKRKTTLAPNT
						QTASPRALADSLMQLARQVSR
						LE/QRAKREPENEEEVDILSLSE
						PVKINIKKEQEEKQEEVKFYLPP
						TPGSEFIGDVTQKRLEEVDSTFG
						DDLVQGFSGGISRRYGEKPGVQ
						DLEVEPDSVTEMLQFYD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22154	52522	A	22282	<u> </u>	1197	MAWPCISRLCCLARRWNQLDR
		` `				SDVAVPLTLHGYSDLDSEEPGT
į						GGAASRRGQPPAGARDSGRDV
						PLTQYQRDFGLWTTPAGPKDPP
						PGRGPGAGGRRGKSSAQSSAPP
						APGARGVYVLPIGDADAAAAV
						TTSYRYGLGRDRNESPPCGGRT
						RRPGARGMGWAAERRRLEPGS
						HATSELPAASEVTPVWSVGTA
						GGAFAAPCPETVLEHPRAGSAP
						LPSQPPSWGQPSEWPAFSRVGT
						GLPLTPTAGPSRARGARRPCPP
						ALPGHCLLDRTYTGLQTLGAET
						LLAVVNSAAMNVGVQVVDVE
						LHRHSLGEDCIYPQSSESDISDA
						PPSLPLTIPAPVKASSPIKQSHEP
						VPDTSVEKGS\PGSCPFHL*GPL
						SHLGSSPGFLLWRPPGLLSSVA
22155	52523	A	22283	131	677	PSSLS/CDIFLRSPISTPSPSPLPRT
22133	32323		22203		077	PTSTPVHVKQGTAGSVINNPYV
						IMDKQPGQVIGATTPSTGSPTN
		ŀ				KISTASQISQGTGSPVPKIHGSSF
1						VTSTVKVIIKQEPGEAPHVPAT
						GAASQSPLPQYVTVKGGHMIA
						VSPQKQVITPGEGIAQSAKVQP
						SKVL/GQIG*CLPTLARADLLYS
22156	52524	A	22284	3	5128	LLKIHSSQSSPQQAVLTIPSQLK
22130	32321	` `	2220 !		3120	PLSVNTSGGVQTILMPVNKVVQ
						SFSTSKPPAILPVAAPTPVVPSS
		İ		İ		APAAVAKGEDASCFSAKSVEQ
				}		YYGWNIGKRRAAEWQRAMTM
						RKVLQEILEKNPRFHHLTPLKT
	•				'	KHIAHWCRCHGYTPPDPESLRN
	1					DGDSIEDVLTQIDSEPDHGSLG
						DGKSSSVGPTAVVTLLACHLEC
ŀ						PSSFSSADNLCRKLEDLQQFQK
						REPENEEEVDILSLSEPVKINIKK
						EQEEKQEEVKFY
22157	52525	Α	22285	112	651	GGLCTMARTKADWPANRTGG
	32323	ļ .				KAP\RKQLAYKKPASQECGPLT
						GGVKKPHRYR\PGTVGGSVKFR
						R\YQKST\ELLIRKLPFQ/RRLVR
						EIAQGL*NKILRFQSAAIG\ALQE
						GKWRPNLVWPFLKDTNLVC*S
						MAKRVTIIAKKTFQLAPPHTVE
						NRALRIHSWMGKHFISQKKKK
						NFSSSCYW
22158	52526	A	22286	1	339	112 5550 1 17
22130	12220	1,,	22200	<u> 1</u>	1227	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22159	52527	Α	22287	2	1369	EPAPPARRSLSAAAAASGTTVD
						TASHSLVFSQQ\LNMQREIGFRV
						NCTVAM*DVYFKAHRAVLAAF
						SNYFKMIFIHQTSECIKIQPTDIQ
						PDIFSYLLHIMYTGKGPKQIVD
						HSRLEEGIRFLHADYLSHIATE
ŀ						MNQVFSPETVQSSNLYGIQISTT
			,			QKTVVKQGLEVKEAPSSNSGN
						RAAVQGDLPQLQLSLAIGLDDG
						TADQQRACPATQALEEHQKPP
						VSIKQERCDPESVISQSHPSPSSE
						VTGPTFTENSVKIHLCHYCGER
						FDSRSNLRQHLHTHVSGSLPFG
						VPASILESNDLGEVHPLNENSE
						ALECRRLSSFIVKENEQQPDHT
						NRGTTEPLQISQVSLISKDTEPV
						ELNCNFSFSRKRKMSCTICGHK
						FPRKSQLLEHMYTHKGKSYRY
						NRCQRFGNALAQRFQPYCDSW
						SDVSLKSSRLSQEHLDLPCALES
						ELTQENVDTILVE
22160	52528	Α	22288	363	947	STARRGKSRSRPAVALQRQPSA
						ASRSPSSSRRRRGVSKEVSVPLA
						RTKQTARKSTN\GK\APRKQLA
					İ	YKKPARKSA\PSTGGVKKPHRY
						RPGYCGRSVKFRR\YQKSH*DF*
				İ		FRKLPFQAVWVRENLLKDF*K
		ĺ				QILRF\QSRSYRVALQGGQVEA
						LSGVGLF*KTTNLLC*SMPKRV
						NNYCQKDIPAKHARIPWRTVL
22161	52529	A	22289	1	92	
22162	52530	С	22290	10	198	
22163	52531	В	22291	74	205	

SEQ ID		Met	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	F	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22164	52532	A	22292	2	1914	RGHCEPSRAAPSRADRHPASAL
						ARDPGRRRGPAGQAAPQPMEP
		ļ				ENIVANTVLLKAREGGGGNRK
						GKSKKWRQMLQFPHISQCEEL
						RLSLERDYHSLCERQPIGRL\LF*
						EFCATRPELSRCIAFQDGMAEY
			!	ļ		EATPDEKRKACGRRLMQNFL\S
						HTGPDLIPEVP\RQLVTNCTQRL
						EQGPCKELSQELTRVTHEYLSV
	<u> </u>					APFADYLDSIYFNRFL\QGKGLE
			l i			REAI\TKNTLRQYRVLGKGGFG
						EVCACQVRATGKMYACKKLE
						KKRIKKRKGEA\MSLNEKQILE
						KVN\SRFVVSLAYG\YETKD\AL
					i	
						CLVLTLMNGGDLKFHIYHMGQ
						AGFPEARAVFYAAEICCGL\EDL
		1				HRERIVYRDLKPENILLDDHGHI
			:			RISDLG\LAVHVPEGQTIKGRVG
						\TVGYMAPECLKNERYTFSPDW
						WALGCLLYEMIAGQSPFQQRK
						KKIKRE\EVERLVKEVPEEYSER
						FSPQARSLCSQLLCKDPAERLG
						CRGGSAREVKEHPLFKKLNFKR
1						LGAGMLEPPF\KPDPQAIYCKD
						VLDIEQFSTVKGVELEPTDQTF
1		l				YQKFAT\GSVPIPGKNEQMVET
		1				ECFQELNVFGLDGSVPPDLDW
						KGQPPAPPKKGLLQRLFSSPKD
						CCGN\CSDSEEELPYPPLAPSPRP
22165	52533	Α	22293	102	403	
22166	52534	A	22294	1086	1436	
22167	52535	Α	22295	965	1231	
22168	52536	Α	22296	1434	2215	TPSFSLRTGVNKEHWVLLGFST
						EDGRW\MFTRGGE\DCTARIWD
						LRSRNLQCQRIFQVNAPINCVC
						LHPN\QAELIVGDQSGAIHIWDL
		İ				KTDHNEQLIPEPEVSITSAHIDP
						DASLHGQLFNSTGNCYVWD\LT
				}		GG\IGDEVTQFIPKTKIPAHTRY
				1		ALQCRFSPDSTLLATCSA*LRR/
			1			CKIWRTSNFSLMTEL\SIKSGNP
				j		GE\SSRG\WMWGCALIGGTPSNI
						VTASSNNLARLWCVKTGEIKRE
						YGGHQKAVVCLPFNDSVLG
22169	52537	В	22297	158	385	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 · · · · · · · · · · · · · · · · · · ·
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22170	52538	A	22298	145	1232	LPVRAEPTRAAAMSGDEMIFDP
						TMSK/NMKKKKKKKPFMLDEEG
						DTQTEETQPSETKEVEPEPTED
						KDLEADEEDTRKKDASDDLDD
			1			LNFFNQKKKKKKK\TKKIFDIDEA
}						\EKGVKDL\KI*SDVQEPTEPED
İ						DLDIMLGHKKKKKKNVKFPDE`
						DEILEKDEALEDEDNKKDDGIS
1						FSNQTGPAWAG\SERDYTYEEL
						LNR\VFNIMREK\NPGYWLPGG
						RKRKFGPWKPPQV\VRVGTKKT
						FFCPLLQDICKLLHRSAQSISFA
						FLLG*IGVPSGSIDG**PNL*SKG
						RF\QQKQIENVLRRYI\KEYVTC\
						HTC\RSPDT\IL\QKDTRLLFPYS
						AETC\HS\RCFCWPVFKTGLPQA
						VHGQSRGTAPVPKA
22171	52539	В	22299	50	319	
22172	52540	Α	22300	3	235	LLLYPLRSLPENPLRLVQQIYSF
						LKGSSEK*NDLPRVPLLELGVG
		1				LLPPNQYSCSRAKPCLTCGLTM
		ļ	22221		2010	SLDGAQQGFW
22173	52541	Α	22301	3	2849	WAKPVARASPAAVPGRQCQIR
		1				GAWSWGEGRGQPGPAPSPAGS PATSEESLAPMAWRCPRMGRV
		1			•	PLAWCLALCGWACMAPKGTQ\
						AEESPFVGNP\GNITGARGLTGT
1						LRCQLQVQGEPPEVHWLRDGQ
						ILELADSTQTQVPLGEDEQDDW
		ŀ				IVVSQLRITSLQLSDTGQYQCLV
						FLGHQTFVSQPGYVGLEGLPYF
						LEEPEDRTVAANTPFNLSCOAO
						GPPEPVDLLWLQDAVPLATAP
						GHGPQRSLHVPGLNKTS
22174	52542	A	22302	3364	3899	LVFLFDDLGIVELVLPGNIVHGI
	32312	1	22302			MSVRPRVAAGLRVRLQPGRRW
		1				LRGPLQLGAPLGLDLLVDVPGS
				İ		PRFVVGVELPLVV/EAPRPPTPP
						AAAQQARS*GLECPGRGAAAP
						REV\LHLQTAHIHGPGPRGGPR
						ARPGLPGLCPP*T/FSCQEAKGE
						RGGAHSNGGSGSKWPPPPPPPS
	<u></u>			L	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22175	52543	Α	22303	1660	4652	PNPNTCLCNHVESGPCCPSTHT
						YTLTDLQPLFGVRVPTRPSGRG
						TQAEESPFVGNPGNITGARGLT
						GTLRCQLQVQGEPPEVHWLRD
						GQILELADSTQTQVPLGEDEQD
						DWIVVSQLRITSLQLSDTGQYQ
						CLVFLGHQTFVSQPGYVGLEGL
						PYFLEEPEDRTVAANTPFNLSC
						QAQGPPEPVDLLWLQDAVPLA
						TAPGHGPQRSLHVPGAAVGAW
		ł				EGGSRLWLRAPEAGRKRWGGE
						VPPVTHAGAGMLRSERD
22176	52544	Α	22304	54	288	SEAGQAQPAAAVGFVWTGRRA
		-				WETGEGYYSNSLSMELQL*RQ
		1	}			CI*DCFFSSSWRCSVPVYFKEEK
						YKGNLVCLLFFK
22177	52545	В	22305	243	2641	
22178	52546	Α	22306	35	204	
22179	52547	Α	22307	244	424	PSLFKNIMTFYPNLHFLWCRGR
		1		}		LYLDVWRQ/VR*FVRNTNYQLP
		1				LVLHNLLSRFNTFDELV
22180	52548	Α	22308	1	477	
22181	52549	Α	22309	2	418	WYQHLLLMRASGSFQSWWKV
						KGEPAYHMANRP**SLISLRTK
						AKTGA*LFDYFVPHFIRKN*SLF
						TIAKAWNHPKCTSVTDWIKKM
					1	WYIYTMEYYAVTRRNKIVSFA
					·	ET*MELEVIIVSKLTQEQKTKHC
						MFSLTSGS
22182	52550	Α	22310	30	199	
22183	52551	A	22311	547	682	LVFLTNQRLVSTHQGTW\WE*K
						*CQPVIPSL*FYFLRKFSVIVIW*
						*LLILP*HPMALKI/C/CCQAACS
						GHFTAGTGEITRPCSISSKVLKR
						LSRL*STSGS**LVFLTNQRLVS
		1				THQGTWMKLEAIILSKVTQEQK
						TKHHMLSLRSGS
22184	52552	Α	22312	2	194	CPSMIDWLKKMWHRYTMEYY
						AA\IKKNEIMS\FAGTWMKLEAI
						ILSKLTQEQKN\KHCMFSLITGS
22185	52553	С	22313	1	1845	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22186	52554	A	22314	1	1384	MRFQDRTWQVPYPSQCSWGSS
						RQGYVPQARGIVLDSQEAYPSP
			ļ		İ	GGTTLCMHDPDKQAPG/PVRRP
					ì	APGN\PNRLVVSVALQFLTFHES
						GITSALRVRPLPQSV\SFQGPGC
						TEA*RRASICRRRVS*HSTKLDL
						GAVAAQQQAWVAG\PRSPQRR
		1				CHQYS*AGPAWLGLQRGRGSH
		1				RPDTSPWHWFWALAGREEQVL
		ŀ				PPESLL/RARTPGTHLAPGPGA*
					1	TRVQSGPGSWLTWAG*GTPQVI
						SS/PGAGAVGAPETTAVPTVSIA
						PRRSVEMWPETKEHLVAVARR
						KWVSAQEKRLSLRGCGSEKMG
						VSPGETRVYFPALTS\RCRRRCG
ŀ		İ				RVSEPSPTASWAPAPSLGWAYQ
	,					LAGGRLVLRVGLSAGWRAACP
		ĺ				QGGPISWLEGGLSSGVGLSAG
						WRAACPQGGPISWLEGGLSSST
ł		ĺ				MRQPRAPQSSLPAPAPWGAPDP
						LAMLAAWIEARPGVWPTQPGP
						WQVSWVCV
22187	52555	Α	22315	11	431	RHLLAAPPPAPPAAAAAAALSH
						CLRSSGRLAPHTSRRLPRVVKR
						RVNALKNLQVKCAQIEAKFYE
						EVHDLERKYAVLYQPLFDKRF
		t				EIINAIYEPTEEECEWKPDEEDEI
						SEELKEKAKIVDEKK\DEEKEDP
						KGIPEFW
22188	52556	Α	22316	64	467	PAAWLPILVAARQLTVQMMQN
						PQILAALQERLDGLVETPTGYIE
						SLPRVVKRRVNALKNLQVKCA
				1		QIEAKFY\EEVHDLQRKYAVLY
						QPLFDKRFEIINAIYEPTEEECE
						WKPDEEDEISEELKEKAKIEDE
22189	52557	Α	22317	2	177	
22190	52558	В	22318	228	377	
22191	52559	Α	22319	630	767	WRVKFKNTVKIQANF*KSRFIH
				1		LTVVPVTILSQVTLQLTMSPKT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22192	52560	A	22320	80	1362	VTACAAPAAWLPILVADIWSSY
						NMADIDNKEQSELDQDLDDVE
1						EVEEEETGEETKLKARSA*LFR
1						MM\QNPQILA\ALQERLDG/LW
						VETPT\GYIESLP\RVVKRR\VNA
						LKNLQVKCAQKETQ\FYE\EVH
Ì						DLERKYAVLYQPLFDKRFEIIN
ŀ						AIYEPTEEECEWKPDEEDEISEE
1	ļ					LK\EKGQIEDEKKDEEKEDPKGI
						P*IW\LTVFK\NVDLLSDMVQEH
					j	DEPILKHLKDIKVKFSDAGQPM
İ						SFVLEFHFEPNE\YFTN\EVLTKT
						YRMRLEPEPDDSDHFSFDGQEI
						MGCTGCQIDWKKGKNVTLKTI
						RK*/RRNHKGRGT\VRTGH*NQ
						FPNDSFSNFFCPLLKSPESGDLD
						ED\AEAILAADFEIG\HFL\RERII
						PRSVLYFTGEAIEDDDDDDYDEE
	ŀ		ļ			GEEADEEGEEGDEENDPDY\D
						PKK\DQNPAECKQQ
22193	52561	Α	22321	3	425	GCGCGGAAARGSPAPGRPAAS
				ľ		PPDCPLWLQGCYPPPSCDALSA
				ļ		TEADGCD*IYPPDTSRDGPHQA
						SLPGESSSFPGQPNDSYLPERGS
						ERSRQASYATPAAETQDPDEGL
						PQPGPEALPGGLQVPKSAAPFC
						GAQHAAGQ
22194	52562	Α	22322	2	285	SRFEPRVRPTGGDRPHQASPPG
						DSSSFPGQPNDSRLPECGSECRG
						QASYATPAAETQDPE*RSSPTRS
						*SP\PGGFQVPKSAAPFCGAQHA
						AGQ
22195	52563	Α	22323	85	473	FTMPAVALVPKELYLSSSLKDL
						NKKTEVKPEKISTK/QDYVHSIL
						GTANIKKAIEEAERLSESLKLRY
						KEAEVRK*LEEKDRQKEAQQL
						QQKRQETGRE\DGGTLAKVSLE
L						NVLDYQDKTQKSNGEKSEKM

	SEQ ID NO.	INICE	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22196	52564	Α	22324	1	1449	MMRAHMFVYKELKQIYKKKT
						HPHQKVGKGYKQTLLRGRHLR
						GQETYEKKLTHVYETTDFKQQ
						QYYFHSILGPANIKKATGETER
<u> </u>		l				LSESLKLRYEEVEIWKKLEEKD
						RQGEAQWLQQKRQETGREDGS
						MLAKGSLEIVLDSKDKTQKSN
			:			GEKNEKCETKEKGAITAKELYT
						MMMDKNISLIIMDAQRMQDYQ
						DSCILHSLSVPEKAISPGVTASW
		l				IEAHLPDDSIDTWKKRGNVEY
[MVLLDWFSSAKDLQIGTTLWH
			,			LKDALFKWEKGGYKNWFLCY
						SQYTTNAKVTPPPQHQNEELSIS
			;			LDFTYPSLEESIPSKPAAEMPPP
			i			PIKVDEDIELISDQISDNDQNER
						TGPLNISIPVESVAASKSDVSPII
						QPVPSIKNVPQIDHTKKLAVKL
		İ				PEEHIIKSESTNHEQQSPQNEKV
						IPDCSTKPVVSSPTLMLTDEEKA
						HIHAETALLMEKNKQEKELQE
						RQQGKQ/K/EKLRREEHEQKAK
22197	52565	Α	22325	185	2983	FIMPAVASVPKELYLSSSLKDL
						NKKTEVKPEKISTKSYVHSALK
		İ				IFKTAEECRLDRDEERAYVLYM
						KYVTVYNLIKKRPDFKQQQAD
						CAYFLGQTVHILFQHRALLEKK
			•			YRAVGAITAKELYTMMTDKNI
		l	,			SLIIMDARRMQDYQDSCILHSL
		l				SVPEEAISPGVTASWIEAHLPDD
						SKDTWKKRGNVEYVVLLDWF
		ŀ				SSAKDLQIGTTLRSLKDALFKW
		l				ESKTVLRNEPLVLEGGYENWL
						LCYPQYTTNAKVTPPPR
22198	52566	Α	22326	313	3720	FIMPAVASVPKELYLSSSLKDL
		l				NKK\TEVKPEKISTKSYVHS/AP
						RKIFKTAEECRLDRDEERAYVL
						YMKYVTVYNLIKKRPDFKQQQ
						DYFHSILGPGNIKKAVEEAERLS
						ESLKLRYEEAEVRKKLEEKDRQ
						EEAQRLQQKRQETGREDGGTL
						AKGSLENVLDSKDKTQKSNGE
		•				KNEKCETKEKGAITAKELYTM
						MTDKNISLIIMDARRMQDYQDS
						CILHSLSVPEEAISPGVTASWIE
						AHLPDDSKDTWKKR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22199	52567	A	22327	46	562	LLQGPGFCGDSMPVEARPSAA
						WRGGRAPGWTPFSPPDRETWK
Ì						PQ/RASQGESGQRRFGPCPRSGG
						RPPGPGSAAHSSRSQRWRKASR
						AWPAAPRKGLLPPAQAWPPTP
		İ				AAGL*GFLAACKR/PPVGGRGP
						APAPCRPASSRCFLICDSGPPCL
			! !			CSASRAAWMGSAASSRVLFIEG
22200	52568	В	22328	51	445	
22201	52569	Α	22329	51	366	CQWNTQTQVCYKQDVQSITAL
						GSAAQCTCLSSPR*SGKFKLLFS
		1				PGLPGTLVPSTIPATSAPRA/SHR
						AGEAWVGEPGRALAALQNSS\P
						DWAGYQGLPCKPSIRAA
22202	52570	Α	22330	257	916	SPSGSNLETHLRAPDLGGPCPP
						GPRWWA*TPQPGHPQTRPSPAA
	,					QPCPFPPQCQQLIHCPC/SPAPGT
						VAAPPAP\PTPASQCLLCPWTSA
						PVCSPAPTFPCPMAPVMAFMLP
		l				SYSFP/PGTPNLPQAFFPSQPQFP
		İ				SHPTLTSEMASASQP\GSPAGPR
		1				SPDSHVLVQPPGPPHHRPWVGP
						PHRSFSPAAARPCSSTCCSWRK
						PLRVALEPWGPQGPQRQQL
22203	52571	Α	22331	151	1232	GRLGDARGPQGPEAPGGEEAA
						RADSTRACRSCGLLLGGRRGCR
						PSLKNAELLELTVRRVQCVLRG
į						RAREREQLQAEARRALRCRLHP
						VACTKVHTFVSTCQAIDATVA
						AELLNHLLESMPLREGSSFQDL
						LGDALAGPPRAPGRSGWPAGG
ŀ		ŀ				APGSPIPSPPGPGDDLCSDLEEA
						PEAELSQA/PC*GARLGARSPGQ
						PDHSPNCPGVSGGLGDQCQPES
ŀ						CRGGARPSLGSPPLLPGVQMW
						WG*GPGSLPGSSLPPLMDGLQG
						QPLVTSPAQRPSPVS*KNF*GPC
1						SSGSGWREGATGRRKNFVELP
						ALSQVSPSPAFNPRLCGLWGQR
						WQKLVLGTSVPGSPGLNKSLN
						LPLQRGDERQVH
22204	52572	Α	22332	1	1524	

sequence 09/540,217 codon for peptide sequence deletion	codon, /=possible nucleotide n, \=possible nucleotide insertion) SLLQEVGIICLARGPGCGPE
22205 52573 A 22333 3 8875 EDGS	SLLOEVGUCL ARGPGCGPE
	SLLOEVGUCL ARGRACGRE
	SEEQE VOIICEAROI GCGI E
	SPATGLALQLQGPAAMG
	SWWILWAGATLLW\GKSG
	ISGPTTVFSQPGLTQEASV
1 1 1 1 1	NTGREEFLTAFLQNYQLA
	AYPRLLISSLSESPASVSILS
1 1 1 1 1 1 1	NTSKKVTVRPGESVMVNI
	AEMIGSKIFQHAVVIHSDY
:	QALNAKPDTAELTLLRPIQ
	TEYFVLTPPGTSARNVKEF
	AGAAGASVSVTLKGSVTF
1	FYPAGDVLRVT
	DGPLAPCHGLVPPAQYFQ
	DACQVQGHPGGLCPAVA
, <u> </u>	AACQAAGAQLGEWRRPDF
	CPAHSHYELCGDSCPVSC
PSLS	APEGCESACREGCVCDAG
FVLS	GDTCVPVGQCGCLHDGR
YYPI	GEVFYPGPECERRCECGP
GGH'	VTCQEGAACGPHEECRLE
DGVC	QACHATGCGRCLANGGIH
YITL:	DGRVYDLHGSCSYVLAQ
VCHI	PKPGDEDFSIVLEKNAAGD
LQRL	LVTVAGQVVSLAQGQQV
TVDC	GEAVALPVAVGRVRVTAE
GRNI	MVLQTTKGLRLLFDGDAH
LLMS	SIPSPFRGRLCGLCGNFNG
	DDFVLPNGSAASSVETFG
AAW	RAPGSSKGCGEGCGPQGC
PVCL	AEETAPYESNEACGQLRN
PQGP	PFATCQAVLSPSEYFRQCV
	CAQKGDKAFLCRSLAAYT
AACC	QAAGVAVKPWRTDSFCPL
	AHSHYSICTRTCQGSCAAL
SGLT	GCTTRCFEGCECDDRFLL
SQGV	CIPVQDCGCTHNGRYLPV
NSSL NSSL	LTSDCSERCSCSSSSGLTC
1 1 1 1 1 1 1	GCP\PG\RVCEVKAEAR\N
1 1 1 1 1 1	TRGL\CVLSVG\ANL\TTF
1 1 1 1 1	RGATTSPGVYELSSRCPGL
1 1 1 1 1 1	PWYRVVAEVQICHGKTE
	QVHIFFQDGMVTLTPNKG
	NGLRVDLPAEKLASVSVS
	OGSLLVRQKAGVQVWLGA

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22207	52575	A	22335	3	7639	ITCPENSHYEVCGPPCPASCPSP
22207	32373		22333		7037	APLTTPAVCEGPCVEGCQCDA
						GFVLSADRCVPLNNGCGCWAN
						}
						GTYHEAGSEFWADGTCSQWCR CGPGGGSLVCTPASCGLGEVCG
1						LLPSGQHGCQPVSTAECQAWG
1						DPHYVTLDGHRFDFQGTCEYL
						LSAPCHGPPLGAENFTVTVANE
						HRGSQAVSYTRSVTLQIYNHSL
1						TLSARWPRKLQVDGVFVTLPF
						QLDSLLHAHLSGADVVVTTTS
22200	50.556	ļ. —	100000		224	GLSLAFDGDSFVRLRVPAA
22208	52576	Α	22336	2	224	HCYMVVQLWKPRSLRRKSDQL
1						WSSWNQLNTSELSSFISAEMQT
						EAFDCGSMAGRM*GQGRYLLA
ŀ						ALWDSSVPLS*PQAEPWSGVGP
						AHLTE/LLSPSRFVSISDLLAPKD
						LGTESQVSMEV*KDMGRFHWE
						DIPP*A*VFVIFVFQIFISRTY\DA
						TTHFETTCDDIKNLL*R*QGRYL
		 			1006	LAALWDSSVPLS
22209	52577	Α	22337	2	206	LSSSYLDQPCYETINRIKLYSES
						LARYGKSPYLYPLYGLGELPQG
22210	50.570	ļ	22228	12	457	FAR*EPVFKTSVIQNLIFNIMCI
22210 22211	52578 52579	A	22338	2	457 1556	FVFPSEVAVGLALSPASIFLSFSS
22211	32379	A	22339	2	1330	PFLRAQERLPSLLWWARSPCAF
		l				SLLGSCVRACPAMNEEYDVIV\
						LGTG\LTECILSGIMSVNGKKVL
						HM\DRNPYYGGESASITALEDL
		1				YKRFKIPGSPPRVKWGEERDW
						NVDL\IPKFLKANG\OLVKMLL
						YTEVTRYLDFKVTEGSFVYKG
						GKIYKVPSTEAEALASSLMGLF
						EK\RRFKEFLVYVANFD\EKDPR TF\EGIDPKKTTMRDVYKKFDL
						GQDVIDFTGHALA\LYRTDDYL\
						l [~]
						DQPCYETINRIKLYSESLARYG
						KSPYLYPLYGLGELPQGFARLR
						CYLWEGTYMLNKPIE\EIIVQE
						WGKVIGCKI*REEIARCMQLIC
						DPSYVKDRVEKVGQVITVICIP
		1				QPPPSKNTNDANSC\QIIIPQNQ\
						VNRKSDIYVC\MISFA\HNVAAQ
		1		,		GKSIAIVSTTVET\KEPEKEIRPA
						L\ELL\EPIGTEILVSISDLL\VPKD
		1				LG\TESQIFIS\RTYDATTHFETTS
20215	50.500	_	222.40		100	D\DIKNIYKR\MTGSEFDFEEMK
22212	52580	A	22340	66	420	
22213 22214	52581 52582	A	22341	2	140 254	
22214	32302	A_	22342		434	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22215	52583	Α	22343	82	3530	HLIVAMPEPTKKEENEVPAPAP
						PPEEPSKEKEAGTTPAKDWTLV
						ETPPGEEQAKQNANSQLSILFIE
						KPQGGTVKVGEDITFIAKVKAE
						DLSEKPTI/KWF/K/GKWMDLAS
						KAGKHLQLKETFERHSRVYTFE
						MQIIKAKDNFAGNYRCEVTYK
						DKFDSCSFDLEVHESTGTTPNID
						IRSAFKRSGEGQEDAGELDFSG
						LLKRREVKQQEEEPQVDVWEL
						LKNTKPSEYEKIAFQY/GITDL\R
						GMLKRLKRSI
22216	52584	Α	22344	66	358	TWVRGNPKPKITWMKNKVAIV
						DDPRYRMFSNQGVCTLEIRKPS
						PYDGGTYCCKAVNDLGTVEIE
						CKL\EVKVIYQGVNTPGQPVFL
		<u> </u>				EGQQQSLHNKDF
22217	52585	Α	22345	67	3515	HLIVAMPEPTKKEENEVPAPAP
						PPEEPSKEKEAGTTPAKDWTLV
						ETPPGEEQAKQNANSQLSILFIE
						KPQGGTVKVGEDITFIAKVKAE
						DLLRKPTIKWFKGKWMDLASK
						AGKHLQLKETFERHSRVYTFE
						MQIIKAKDNFAGNYRCEVTYK
						DKFDSCSFDLEVHESTGTTPNID
						IRSAFKRSGEGQEDAGELDFSG
						LLKRREVKQQEEEPQVDVW\EL
						LKNAKPSEYEKIAFQYGITDLR
22210	52596		22246		20	GMLKRLKRMRREEK
22218	52586	Α	22346	2	39	FQFCVSLFVYIQLNSDVLPE*PR HKKKPPTLSHWFLK*NLHL*N*
	:				:	NTKALKYRKLV*LTSQKH*QM
						TYPFCNYLKTN*VTTNTKVK*T
						EGY*TAKTQKLYPDKVVMATN
						KPLKTRSPK\PAKYGSPHDTAT
						WTGSGLGLEAS*HSA
22219	52587	A	22347	1	113	W I GOOLGELAS TISA
22220	52588	C	22348	47	337	
22221	52589	Ā	22349	1	2292	
22222	52590	A	22350	2	441	LRHAKWFQARANGLKSCVIVIR
						VLRDL\VTRVPTWGPLRGWPLE
						LLCEKSIGTANRPMGAGEALRR
						VLECLASGIVMP/QHALRLAAF
						GQLHKVLGMDPLPSKMPKKPK
						NENPVDYTVQIPPSTTYAITPM
						KRPMEEDGEEKSPSKKK
	<u> </u>		<u> </u>	I	<u> </u>	THE PRODUCTION OF STATE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22223	52591	A	22351	45	402	HALAALEKLFPDTPLALDANK
22223	32391	A	22331	43	402	KKRAP\VPVRGGPKFAAKPHNP
		1				GFGMGGPMHNEVPPPPNLRGR
						GRGGSIRGRGRGRGFGGANHG
						GYMNAGAGYGSYGYGGNSAT
00004	50.500	ļ. —	222.52	242	2112	QGYK*LFHKTAYGYS
22224	52592	Α	22352	243	3113	FPDTTEEVKMRPMRIFVNDDRH
						VMAKHSSVYPTQEELEAVQNM
						VSHTERALKAVSDWIDEQEKG
1						SSEQAESDNMDVPPEDDSKEG
						AGEQKTEHMTRTLRG\VMR\VG
						LVAK\CLLLKGDL\DL\ELVLLC\
						KEKPTT\ALL\DKVADNLAI\QL
						AAVTEDKFEILQSVDDAAIVIK\
						NTKEPPLS\LTIHLTSPVVREE\M
						EKVLAGETLS\VNGPPGNVLDR
						SRIAFAAFRGSLPETQRWFQ\AR
						ANGLK\SCV
22225	52593	Α	22353	139	245	
22226	52594	Α	22354	875	2019	NPNSQMPKMTRSTKQLTNKSG
						QIDGCPGCRVVPTWCRLGHLSH
		ļ				YWAVWASILRFWEVWKYWLG
1						NMGRKSSEFPILKLNG\RTMGD
						*VLPFHMYV\QPVV\HPQLYPHS
				:		VTELTGDYFKPWWDGQPSLQQ
						VL\ERVDEWM\AKEGLL\DPNV
			ŀ			QVQFLFTLWEDWGLKKSVLPG\
						QCQ*LGAWPVGGFTSKQWD*S
			1			*KKAYSF\AMGCWPKNGLLDM
			•			NKGLSLQHIGRPHSGIDDCKNI
1						ANIMKTLAYGGFIFKQTSKPVLI
						GRGODGAGOGSLVGPPONPPL
						PHQRGKRESGHKLCCPKVVPSA
						SLWGSMPLGRGLCCPLQGALA
						TWPFLEARHFCAPPSPTPQSQPS
						ISPSHCGPGLRGTPGTHRLLPW
						YNRLLWGHQDPRCAPSHP
22227	52595	Α	22355	507	916	RKKNEKEGTKPPRNMGICEKT
					-	KPTFDWCT*K*RGEWNQVGKY
						TSGYYTGELPQSSKTGQHSNSG
						NTENTTKILLEKSNPKTHNW*I
						HQG*NEGKTVKGSQREK*GYP
						QREAHQTNSRSLCRNPTRQKRV
22228	52506	D	22256	72	250	GVNIKHS
22228	52596	В	22356	72	359	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22229	52597	Α	22357	1236	1987	GNSSPPGLPCKSSSSPPPPPSLTG
		İ				APARRAAAAPPRPASRPRPPLA\
					ł	GGGVPQLFATRPP/PGPTRARRP
1	1	1				TWR*GPRGSPEETPRRRREWTR
						RPEPGCRAQGSRRPARPPSGSS
						AVRSASPASSPAPAASAPLGPRS
						RWS*RPPSRPRR/SPPPPPPGRSR
						SDRER/GRRKEASPPLTLAEAPA
					ļ	Y/RQDQAPPPGSPHLGFQSAAG
						PSRTASSAP/RSGIGGRGANTPA
						PAAIFPPSLRPPRAGPVGDLRAS
]	GVDAAA
22230	52598	Α	22358	287	2082	LKTETSEEKARRYKDRPSQLNA
	2000					VFQEQKKMIQAQESITLEDVAV
						DFTWEEWQLLGAAQKDLYRD
						VMLENYSNLVAVGYQASKPDA
						LFKLEQGEQLWTIEDGIHSGAC
Ì	:					SDIWKVDHVLERLQSESLVNRR
						KPCHEHDAFENIVHCSKSQFLL
						GQNHDIFDLRGKSLKSNLTLVN
						QSKGYEIKNSVEFTGNGDSFLH
						ANHERLHTAIKFPASQKLISTKS
						QFISPKHQKTRKLEKHHVCSEC
						GKAFIKKSWLTDHQVMHTGEK
			•			PHRCSLCEKAFSRKFMLTEHQR
						THTGEKPYECPECGKAFLKKSR
			:			LNIHQKTHTGEKPYICSECGKG
						FIQKGNLIVHQRIHTGEKPYICN
						ECGKGFIQKTCLIAHQRFHTGK
						TPFVCSECGKSCSQKSGLIKHQ
						RIHTGEKPFECSECGKAFSTKQ
1						KLIVHQRTHTGERPYGCNECGK
		ŀ				AFAYMSCLVKHKRIHTREKQE/
						CSQGGKSSCREAQLITHQ*CHA
İ						GEKLC*RGDYTSAFCGPSDIIKH
						QRP/HSQNRN/CSPCGTASGQM
						CSLRR*QWICTGQKPCECSECG
						CAFRDQLCLILCYRKPIGRK\PD
						LCGKGLSKNL*FIMWPKSIHCE
22231	52599	С	22359	90	344	LEGROESKIVE THIWI KSHICE
22232	52600	A	22360	2	543	
22233	52601	A	22361	373	490	
22234	52602	A	22362	111	396	GAPPGIWCLSLWACRALSPLSS
22237	32002	l' `	22302	111		AAQHRKRPSALSPATPAGSRPL
						RRCAPGFAATKTQRVQRRPPRR
						*GAAAGAGGS*RPLCGVVIHSV
						LHNTPE
L	l	<u> </u>	L		l	LIMITE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22235	52603	A	22363	1902	3474	VPQRSTGSGRARSARRPLRAPD PCAAAPRVSPQPRPSECSGGRR GGSKTWAKRNAGKELTDRNW DQEDEAEEVGTFSMASEEVLK NRAIKKAKR\RNVGFESDTGGA FKGFKRFGGYPSEEERFSGFG\S GAGGKPLEGL\SNGNNIT\SAPP FASAKAAADP\KVAFGK*LPSPS RLLVDKNVSNPKTNGDSQQPSS SGLASSKACVGNAYHKQLAAL DCSVRDWMVKHANTNSLCDL TPIFK\DYEK\YLANIETANTGN\ SGRNSESESNKVAAETQSPSLF GSTKLQQESTFLFHGNKTEDTP DKKMEVASEKKTDPSSLGATS ASFNFGKKVDSSVLGSLSSVPL TGFSFSPGNSSLFGKDTTQSKPV SSPFPTKPLEGQAEGDSGECKG GDEEENDEPPKVVVTEVKEED AFYSKKCKLFYKKDNEFKEKGI GTLHLKPTANQKTQLLVRADT\ NLGNILLNVLIPP\NMPCTRTGK NNVLIVCVPNPPIDEKNA\TMP\ VTMLI\RVKTSEDADELHKILLE
22236	52604	C	22364	65	283	VIMERAVRISEDADEEHRIELE
22237	52605	A	22365	3	419	
22238	52606	A	22366	1	512	CGRPPWKTGQRSQPSCPPGSSG SGRGIP\PAAWHEGEGPEAVLQ GHRAGRGDPACRGLCCLPSQL GGPTSPTSAASRACGSRGAATL VVKVKWFYHP/CRRPSWARGS ATARMRCTSPATRMRTTCRPSP TSARSWRASSMSRWPGAASAR TGRTSTTGGHLRPHHRAPGDG
22239	52607	A	22367		2940	MRRRRKVAAIHAGPRPRSAAA SSEQRGRGPGAARRAGGAEAA ASGVTGGGGGGGAGQHAGGLG FSAGGRAPSQRPLCGSMPARLF SKQEVDYCGLPAPPASSLRSRG HRQRIYSLNQLLQEALRTGRAR PGEGPGWLVTDKGGLGWGSSH SQAQGPDSQAAWGRAPPAHGS SPAFLQVLDVRPQSSRYLPPGT RVCAYWSQKSRCLYPGNVVRA PPPCFGRLLGVPAASSGVPSTG ASGDEDEDLDSVVVEFDDGD
22240	52608	В	22368	215	356	
22241	52609	A	22369	282	499	YLLHCLPSDPPSFTNSPDGRGRR RRRQRRRRSDPPGRGRTKSSTR VPGSNRFV*FHQR*GRLKEWM ANQDTN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22242	52610	A	22370	105	619	ACQASSHSGPVWGPGERTTPG A*ASARREPSRNQHPNLWVTE Q/I*IASPRTVANPLVPARTPTGP RTNKPKHLIKQGRGQPRQPLSG STGGAGESPASGCAGGQPAHR VWIAPAARGCGQHTPWKAPAG NGQSPRPASATRMIGPSPSLSPP APPAVVGVAALGSRNRCIL
22243	52611	A	22371	1	441	
22244	52612	Α	22372	10	621	LRQLERGLAKGQPSTPVDRPLG PDPSAPGTLAGAALPPLEPPAPC LCQDPQEDSVEDEEPPGSLGLP PPQAGVQPAAAAVSGTTQPLG TGPRVSLSPHSPLLSPKVASMD AKDLALQILPPCQVPPPSGPQSP AGPQGLSAPEQQEDEDSLEEDS PRALGSGQHSDSHGESSAELDE QDILAPQTVQCPAQAPAGGSEE T*LRQLERGLAKGQPSTPVDRP LGPDPSAPGTLAGAALPPLEPP APCLCQDPQEDSVEDEEPPGSL GLPPPQAGVQPAAAAVSGTTQP LGTGPRVSLSPHSPLLSPKVAS MDAKDLALQILPPCQVPPPSGP QSPAGPQGLSAPEQQEDEDSLE EDSPRALGSGQHSDSHGESSAE LDEQDILAPQTVQCPAQAPAGG SEETIAKA

2890

SEQ ID	_		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22245	52613	Α	22373	11	2271	MPGELLAPSCCCRGGPTRDRTD
						LSCDAAAATTILGGDRREPCAL
		ĺ				TPGPSHLALTFLPSKPGARPQPE
		1				GASWDAGPG\LSTEAPLPATLE
٠		1				PRIVMGEETCQALLSPRAARTA
						LRDQEGWGL/SPQGSMVDEREL
						HPAGTPEPPSSESSLSADSSSSW
	1					GQEGHFFDLDFLANDPMIPAAL
	1	İ				LPFQGSLIFQVEAVEVTPLSPEE
:						EEEEAVADPDPDSASYAEADDE
						RLYSGEPHAQATLLQDSVQKTE
	-	ŀ				EESGGLRYDTASAPTRHRSLVS
						PKACSCSHDCAP\AG*RGPHLTP
						GLRYDTASAPTRHRSLISPKAC
						GCSHDCVPAG*RGPHLTPGLRY
						DTASAPTRHRSLVSPKACSCSH
						ACVPAG*RGPHLTPGLRYDSTS
						ASARHRPHLRSRASGCGHPSNL
						AGRSRLCPRDRACGHHGSAGS/
						SAWTGSMSGSA\RHR*WG*A*L
	1	1				TPKGDPGG*EPAGRRPEG\TGT
						GTWTQPQGGLHGCQRPGLADL
						AALPSASSLWAPEPSWPS/SALS
						PRAARG*GQPGGRLASGPGLGP
						AFG*PRGVISRAGRAGHLGSSD
						RAVSSPGAQSREEGPSRRQ*GD
						HRQSQAESQ*EEGPKGNVKAG
						LAADSGSHQDHHPEVQEHPLC
						HRQA*CLQGDPASAHLCG\FWR
•						GQ\T*GP\CPASAQSRS*EV*GAL
						R\DLSLGP*VST\QGPG*A*VQG
[l				KGKG\GGGKGGPGGAGTA*HL
22246	52614	Α	22374	2	211	
22247	52615	A	22375	43	241	
22248	52616	A	22376	184	332	
22249	52617	A	22377	1292	1923	KAGARGNQEGTMASSSG\TQK
						VLTIPRGCYQ*K*SKETLP*CSG
						WPNDAR\EAVVNC\CTEFI\HLIS
						SEANEICNKSEKKTISPEHVIQA
						LESLGFGSYISEVKEVLQECKT
						VALK\RRKASSRFGKPLGIPEEE\
						LFGDQANKELFGKSLDQQQAE\
						LAQQ\EW\LQMQQAA\HQA\QL
						AAASAKCI*SRRGSFSGMEEDG
						GWISEIPPAEFLFLL
22250	52618	В	22378	1	1701	
22251	52619	A	22379	1	147	
	1	1		I	L	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22252	52620	Α	22380	1	484	IRHEGRGQRGKMELVQ\VLKRG
						LQQITGHGG/LFRGYLRVFF\RT
						NDAKVGTLVGEDKYGNKYYE
		l				DNKQFFGRHRWVVYTTEMNG
						KNTFWDVDGSMVPPEWHRWL
						HSMTDDPPTTKPLTA\RKFHLG
						RNH*ISTWTGAPQEQYVPYSTT
		l				RK\KIQEWIPPSTPYK
22253	52621	Α	22381	3	258	DMAAQIAEGMAFIEERNYIHRD
		l				LRAANILVSDTLSCKIADFGLA
		1				RLIED\FTIKSDVWSFGILLTEIV
		l				THGRIPYPGMTNPEVIQN
22254	52622	Α	22382	1	738	
22255	52623	A	22383	233	1898	GAGPTGYLRRLPGWAGTMGC
						GCSSHPEDDWMENIDVCENCH
						YPIVPLDGKGTLLIRNGSEVRDP
		1				LVTYEGSNPPASPLQDNLVI\AL
		ŀ				HSYEPSHDGDLGFEKGEQLRIL
:		ļ				EQSV\EWWKAQSLTTGQEGFIP
;						FNFVAKANSLEPEPWFFKNLSR
						KDAERQLLAPGNTHGSFLIRES
						ESTAGSFSLSVRDFDQNQGEVV
						KHYKIRNLDNGGFYISPRITFPG
					ł	LHELASAITPIASDGLCTR\LSRP
						CQTQKPQKPQKPWWEDEWEV
						PRETLKLVERLGAGQFG\RCGW
	}	ļ				GTTTGTTKVAVKSLKQGSMSA
					<u> </u>	GRLP\AEANLMKQLQ\HQRL\VR
						LYAVVT\QEPIYIITEYMENGSL
						VDFLKTPSGIKLTINKLLDMAA
						QIAEGMAFIEERNYIHRDLRAA
						NILVSDTLSCKIADFGLARLIED
				ļ		NEYTAREGAKFPIKWTAPEAI*S
						RGNSPPRSDVWSF\GDLLPEIFT\
						HGRIP/YPQGIPNPELIQKLERGY
						RMVRP\SNCPEKLYQL\MRLCW
						KERPED\RPTFDYLRSVLEDFFK
						RHRRASNQPQALREPLRKPLCF
22256	52624	Α	22384	3	203	
22257	52625	Α	22385	2	467	
22258	52626	Α	22386	1	468	
22259	52627	Α	22387	1	645	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ŀ				sequence	:	
22260	52628	A	22388	821	2637	AAASAWRATHSTTAAGRRTAL
1						FRSPEPGNTRDTRRRRRATAPD
ĺ		1				CCRPATSPRALDWTSLHTRRHP
Í						VQNYLQWSYWQVLVSRWIAF
						AGLAMADYLISGGTGYVPEDG
						LTAQQLFASADDLTYNDFLILP
						GFIDFIADEVDLTSALTRKITLK
						TPLISSPMDTVTEADMAIAMAL
				:		MGGIGFIHHNCTPEFQANEVRK
				1		VKNFEQGFITDPVVLSPSHTVG\
		ŀ				DVLEGKMRHGFSG\IPITETGTM
		İ				GSKLVGIVTSRDIDFLAEKDHT
						TLLSEVMTPRIELVVAPAGVTL
						KEANEILQRSKKGKLPIVNDCD
ĺ						ELVAHARTDLKKSRDYPLASK
						DSQPQLLCGAAVDTREEDKYR
1						LDLLTQAGVDVIVLDSSQGNSV
		ļ				YQY\ARVLYFKQKYPHLQV\LG
						GNVVTAAQAKNLIDAGVDGLR
		1				VGMGCGSICITQ\EVMACGRPQ
		1				GTCCVTR*AEYARRFGVPIIAD
						GGIQTVGHVVKALALGASTVM
		l				MGSLLAATTEAPGEYFFSDGVR
ļ						LKKYRGMGSLDAMEKSSSSQK
		1				RYFSEGDKVKIAQGVSGSIQDK
		1				GSIQKFVPYLIAGIQHGCQDIGA
		1				RSLSVLRSMMYSGELKFEKRT
						MSPQIEGGVHGLHSYEKRLY
22261	52629	Α	22389	2	332	
22262	52630	Α	22390	3	608	MTAVHAGHLNLKWDPKSLEIR
		1				TLAAERLVTQVTTLVNTNSKGP
		l				SDKKRGHSKKVHVLAASVEQA
		1				AENFLEKEDKIVKESQF\L*E*LL
						AAVEYVRKQ/GKECLRPSLEEH
						LEGIISGAALMGDSSCKHDDHR
						ERIAAECNTVLQ\ACRTCF/SEY
						MGNAGCKERSNSLNSAIDKMT
						KKTRDLRRQLHKAVMDHVSDS
			<u> </u>			FLETNVPLL
22263	52631	Α	22391	2	2864	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22264	52632	A	22392	14	2897	PDWRETKQRPSASGLWNLALA
Ì		l		:		QLAAEMTAVHAGHLNLKWDP
		l				KSLEIRT\LAVERLLEPLVTQVT
						TLVNTNSKGPSDKKRGHSKKV
		l				HVLAASVEQAAENFLEKEDKIV
		l				KESQF\L*E*LLAAVEDVRKQG
						DLMKAAARRRSHDDPCSSVEQ
						GNMVQAA*TLLSAVTCLLILA\
						DMGNVYKLLVQLKVVEDGILK
			:			LRNAGNEQDLGIQYKALKPEV
						DKLNIMAAKRQQELKDVGHRD
		1				QMAAARGILQKNVPILYTAS
22265	52633	Α	22393	1	1584	
22266	52634	A	22394	1	3519	
22267	52635	Α	22395	3	673	
22268	52636	Α	22396	1	1574	DVPRDLEVVAATPTSLLISWDA
]		l				PAVTVRYYRITYGETGGNSPVQ
						EFTVPGSKSTATISGLKPGVDYT
		l		•		ITVYAVTGRGDSPASSKPISINY
						RTEIDKPSQMQVTDVQDNSISV
					}	KWLPSSSPVTGYRVTTTPKNGP
						GPTKTKTAGPDQTEMTIEGLQP
]						TVEYVVSVYAQ\NPSGESQPLV
]						Q\VAVPTIPAPTDLKFTQVTPTS
						LSAQWALPNVQLTGYRVRVTP
						KERTGPMKEINLAPDSSSVVVS
						GLMVATKYEVSVYALKDTLTS
		1				GPAQGVVTTLENVSPPRRARVT
		l				DATETTITISWRTKTETITGFQV
						DAVPANGQT\PIQRTIKPDVRSY
						TITGLQPGTDYKIYLYTLNDNA
						RSSPVVIDASTAIDAPSNLRFLA
						TTPNSLLVSWQPPRARITGYIIK
						YEKPGSPPREVVPRPRPGVTEA
						TITGLEPGTEYTIYVIALKNNQK
						SEPLIGRKKTVQKTPFVTHPGY
						DTGNGIQLPGTSGQQPSVGQQ
						MIFEEHGFRRTTPPTTATPIRHR
						PRPYPPNVGQEALS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>	<u> </u>			
22269	52637	Α	22397	3	2382	QTEMTIEGLQPTVEYVVSVYAQ
			•			NPSGESQPLVQTAVTNIDRPKG
			:			LAFTDVDVDSIKIAWESPQGQV
						SRYRVTYSSPEDGIHELFPAPDG
						EEDTAELQGLRPGSEYTVSVVA
						LHDDMESQPLIGTQSTAIPAPTD
	i i					LKFTQVTPTSLSAQWTPPNVQL
						TGYRVRVTPKEKTGPMKEINLA
						PDSSSVVVSGLMVATKYEVSV
						YALKDTLTSRPAQGVVTTLEN
						VSPPRRARVTDATETTITISWRT
						KTETITGFQVDAVPANGQTPIQ
						RTIKPDVRSYTITGLQPGTDYKI
						YLYTLNDNARSSPVVIDASTAI
			,			DAPSNLRFLATTPNSLLVSWQP
						PRARITGYIIKYEKPGSPPREVV
			•	i		PRPRPGVTEATITGLEPGTEYTI
	:				,	YVIALKNNQKSEPLIGRKKTDE
						LPQLVTLPHPNLHGPEILDVPST
						VQKTPFVTHPGYDTGNGIQLPG
						TSGQQPSVGQQMIFEEHGFRRT
			:			TPPTTATPIRHRPRPYPPNVGEEI
						QIGHIPREDVDYHLYPHGPGLN
			•			PNASTGQEALSQTTISWAPFQD
1						TSEYIISCHPVGTDEEPLQFRVP
						GTS\TSATLTGLTRGATYNIIVE
				ł		ALKDQQRHKVREEVVTVGNSV
						NEGLNQPTDDSCFDPYTVSHYA
ŀ						VGDEWERMSESGFKLLCQCLG
						FGSGHFRCDSSRWCHDNGVNY
						KIGEKWDRQGENGQMMSCTCL
						' '
22270	50.600	<u> </u>	22200	10	7500	GNGKGEFKCDPHEATCYDDGK
22270	52638	A	22398	48	7500	GATMACCAAQGEERVTPCASG
						KRGPAATTSLVLWIPSVPPPVPF
						PTLWPPPSWRRQPPGGIRRDFS
				•		RRLRREANLVATCLPVRASLPH
						RLNMLKGPGPGLLLLAVLCLG
						TAVPSTGASKSKRQAQQMVQP
						QSPVAVSQSKPGCYDNGKHYQ
						INQQWERTYLGNVLVCTCYGG
						SRGFNCESKPEAEETCFDKYTG
						NTYRVGDTYERPKDSMIWDCT
}				1		CIGAGRGRISCTIANRCHEGGQS
<u> </u>						YKIGDTWRRPHETGGYM
22271	52639	A	22399	1	403	NSTNKRRGFCFITFKE\EMEKKY
						HNVGLSKCEIKVAMSKEQYQQ
						QQQWGSRGGFAGRARGRGGG
						PSQNWNQGYSNYWNQGYGNY
						GYNSQGYGGYGYDYTGYNN
						YYGYGDYSNQQSGYGKVSRRG
						GHQNSYKPY
L	L	i	l	<u> </u>		

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22272	52640	C	22400	36	214	
22273	52641	A	22401	3	182	
22274	52642	A	22402	223	452	LESMTIFPYSDDKRAQYIFIQKD
						LLKKPVRHYYIMTLSHWAFNT
		ŀ				RLALQYWGKGHKTQIDSEA*Q
						LRNKTMKANFF
22275	52643	Α	22403	1	1037	MSEEQFGGDGAAAAATAAVG
						GSAGEQEGAMVAATQGAAAA
		ŀ				AGSGAGTGGGTASGGTEGGSA
		ł				ESEGAKIDASKNEEDEGHSNSS
						PRHSEAATAQREEWKMFIGGLS
						WDTTKKDLKDYFSKFGEVVDC
1					,	TLKLDPITGRSRGFGFVLFKESE
		1				SVDKVMDQKEHKLNGKVIDPK
		1		Ì		RAKAMKTKEPVKKIFVGGLSP
				}		DTPEEKIREYFGGFGEVESIELP
		l				MDNKTNKRRGFCFITFKEEEPV
		ĺ				KKIMEKKYHNVGLSKCELK\VA
		1				\MSKEQYQQ\Q\QQWDSRGGCA
						GRA\RGRGGDQAE\GYGKGIQA
		1				RWAIKNSYKPYLNYSICQPYPP
						TAGTSLQALCRADFRFSQARSM
22276	52644	Α	22404	299	1508	HYVGGSSSGGDGAAARAT/AR
						AVGGSAGEQ\EGA\MVAATHG
		l				ARGGVWEADAATGGGTASGG
		1				T\EGGSAE\SEG\AKIYASKNEED
]				EGHSNSSPRHSEAATAQRGRM\
						KMVIGGLSWDT\TKKDLKDYFS
		ļ				KFGEVVDCTLKLDPITG\QSRGF
						GFV\LFKESESV\DKVMDQKEH
						K\LNG\KVNDPKRAKAM\KTKE
		l				PVKKIFVGGLSPDTPE\EKIREYF
						GGFGE\VESIELPMD\NKTN*RG
						RGF\CF\ITFKEEEPVKKI\MEKK
			ļ			YHNVGLSKCEIKVVLLNEQYQ
						QQQQW*SIGGFAGIALVIGGGP
						SQNWNQGYSNYWNQGYGNYG
		1				YSS\QGYG\GYGGYDYTGY\NN
					j	Y\YGYGDYSNQQSGYGKVS\RR
						GGHQ\NSYKPHTKFIPICNLFPN
						RWLKQYFPNLKIQFERGSCHLL
22277	52645	A	22405	3	189	EESEPERKRARTDEVL\PEEAAP
						RRKMRTTRTTCPVSRY\AAP\QL
						LLQKLLQRRRKGSCSRG
22278	52646	В	22406	1	1356	<u> </u>
22279	52647	C	22407	135	260	
22280	52648	A	22408	1	1740	
22281	52649	В	22409	80	371	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide sequence	or peptide sequence	deletion, \=possible nucleotide insertion)
22282	52650	A	22410	257	1200	ISWCCMEQKKRLP\FSKREG\PY
		İ				EFIICPSEDSSPLLRCAFCIGGMS
						VKEQMETIRHPLQTTRGPGPGP
						LFTDGPDPIPEAFLHGLLSIREM
	•					VSTEKVKPVGQESEPRVKPLEL
ļ						DDSASGGVHMMVATPGRLMD
	:	ľ				LLQKKMVSLNICRYLALDEAD
		l				RMIDMGFEGDIRTIFSYFKHTSA
						AAIGLQLALDGGTGSGQKGKC
						EEFPSSMSAVSTGLEASACCLS
						VTMDPLETLSRMASSTSSMSSL
						ARAWLSPEPGDSPGMVGGLPR
		į				ELPGMAPLECGEFQRPESLCAD
		ł				RNELGSQSPEFPGQEEEDEDED
		ŀ				DEEEDEDE
22283	52651	A	22411	107	2069	GGVSPWRACVQQRMEESEPER
		ŀ				KRARTDEV\LPEEAAPRRKMRT
		ŀ				TRTTCPMCRYGSAGSLLLQKLL
						QR\RRKGAAEEEQQDSGSEPRG
		İ				DEDDIPLGPQ\SNVSLLDQHQHL
		ŀ				\KEKAESAAKESCQGRKQLEGK
						KEKILGRIVCRRAE\LDVK*KAD
						WLKGITVLIDPIKTQLGLPPR/Y
						MFLSMSEERHERVRKKYHILVE
						GDGIPPPIKSFKEMKFPAAILRG
						LKKKGIHHPTPIQIQGIPTILSCR
			,			DMIGIAFTGSGKTLVFTLPVIMF
						CLEQEKRLPFSKREGPYGLIICP
						SRELARQTHGILEYYCRLLQED
						SSPLLRC\ALCIGGMSVK\EQ\ME
						TIRQGVHMMVATPG/RALMDL
						LQKKMVSLDICRYLALDEADR
						MIDMGFEGDIRTIFSYFKGQRQ
						TLLFSATMPKKIQNFA\KSALVK
						PVTINVGRAGAASLDVIQEVEY
						VKEEAKMVYLLECLQKTPPPV
						LIFAEKKADVDAIHEYLLLKGV
		ŀ				EAVAIHGGKDQEERTKA\IEAFR
						EGKKDVLVAT\DVASKGLVFPA
		•				IQHVINYDMPEEIENYVHRIGRT
						GR\SGNTGIATTFINKACDESFLI
						D\LKALLVRKPKQKVPPVL\QV
						L\HCGGWSPMLDIGRSSRGCAF
						CGG\LGH\RITDCPKL\EA\MQTK
		<u> </u>				QVSNIGRKDYLAHSSMDF
22284	52652	Α_	22412	435	1428	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22285	52653	A	22413	3	597	HKMAPYSPLVTRLQKALGVRQ YHVASVLCQRAKMAMSHFEPN EYIHYDLLEKNINIVCKRLNWP LTLLEKIVYGHLDDPASQEIEQ GK/S/YLWLWPDHVAMLQFISKI GATTSVFPYNHRMRRYLSK/TG RTDIANLADEFQDHLVPDPGCH YDQLIEINLSELKPHINGPFTPD QAHPVAEVGKVAEKEGWPLDI RVGLT
22286	52654	Α	22414	29	512	
22287	52655	Α	22415	1	543	
22288	52656	В	22416	271	719	
22289	52657	A	22417		2438	SVHKMAPYSLLVTRL*KALGV RQYHVGSVLCQRAR\VAMSHF EPNEYIHYDLLEKSINIVRKRLN RPLTLSEKIVYGHLDDPASQEIE RGKSYLRLRPDRVAMQDA\TA QMAMLQFISSGLSKVAVPSTIH CDHLIEAQVGGEE\DL\RRAKDI NQEVYNFLATAGAKYGVGFW KPGSGIIHQIILENYCVPWLSF*L GPDFHTPNGGG\LGGI\CIGSCG WPNAWEWNGLGIPWELKCPK VIGVKLTGSLSGWPSPKDVILK VAGILTVKGGTGAIVEYQGPGV DSISCTGMATICNMGAEIGATT SVFPYNHRMKKYLSKTGREDIA NLADEFKDHLVPDPGCHYDQLI EINLSELKPHINGPFTPDLAHPV AEVGKVAEKEGWPLDIR\VGL\I GSCTNFKLWKIWGRSAAVAKQ ALAHG\LKCKSQF\TITPGFPRQI\ RATFE\RDGYAQILRDLGG\IVL ANACGPCIGQWDRKDIKKGEK NTIVTSYNRNFTG\RNDANPET HAFVTSPEIVTALAIAGTLKF\N PETDYLTGHGWARSFRLGGSRI AD*VFPKGEF*PRGRDTYQHPP K\DSSGAACWDLSPTSQRLQLL EPFDKWDGKDLEDLQILIKVKG KCTTDHISAAGPWLKFRGHLD NISNNLLIGAINIENGKANSVG\ NPVTQEFGPAPD\LARYYKKHG I\RWVVIGDENYGEG\SSREHAA
22290	52658	A	22418	14	260	RLCIGDAGVSPVCGP/CSAGGGP R*LNLEGGEESSSVPRGG/CCSP VTATTAPGPCFPK*SDGSGCAE APDSGE/CS*ALGPHCGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	1	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22291	52659	A	22419	160	460	RLCIGDAGVSPVCGPLSAGGGP
22291	32039	Α .	22419	100	1400	R*LNL*GGEESSSVPRGG/CCSP
						VTATTAPGPCFPK*SDGSGCAE
						APDSGNCGAFCALRSPHRHPGP
						KPARERLLTGL
22292	52660	A	22420	43	418	SLGCAEQN/FPAIAIQPGMAQEE
	32000	 	22120			RLSRYQQFKDFQRRILVATNLF
ļ						GRGMDIERVNIVFNYDMPEDS
						DTYLHRVARAGRFGTKGLAITF
ŀ						VSDENDAKILNDVQDRFEVNV
						AELPEEIDISTYIEQSR
22293	52661	Α	22421	1	592	RFSKYMPSVKVSVFSRGLSIKK
			-2.2.	•		DEEVLKKNCPHVVVRTRARILA
						LVRNRSFSLQNVKHFVLDECD
						KMLEQLDMRRDVQEIFRLTPHE
						KQCMMF/ERLSRYQQFKDFQRR
,						ILVATNLFGRGMDIERVNIVFN
						YDMPEDSDTYLHRVARAGRFG
						TKGLAITFVSDENDAKILNDVQ
						DRFEVNVAELPEEIDISTYIEQS
22294	52662	Α	22422	2	575	
22295	52663	В	22423	100	412	
22296	52664	Α	22424	2	441	
22297	52665	A	22425	682	1032	
22298	52666	A	22426	l	1324	MAEQDVENDLLDYDEEEEPQA
		ļ				PQESTPAPPKKDIKGSYVSIHSS
						GFRDFLLKPELLRAIVDCGFEHP
						SEVQHECIPQAILGMDVLCQAK
	1					SGMGKTAVFVLATLQQIEPVN
						GQSGQVHIFGLHIGARALVISEH
	ļ				į	ILKDEGAVEGCGVVQCLMARV
						TVLVMCHTRELAFQISKEYERF
						SKYMPSVKVSVFFGGLSIKKDE
						EVLKKNCPHVVVGTPGRILALV
						RNRSFSLKNVKHFVLDECDKM
	1					LEQLDMRR\DVQEIFRLDTTPR
	1					KQCMMFSATL\SKDI\RPV\CRK
						FMQDPMEVF\VDDETKLTLHG\
						LQQYYVKLKTSEKNRSLFDLL\
						DV\LEFNQV\IIFVKSVQRCMAL
						AQLL\VEQNFPAI\AIHRGMAQE
						ERLSRYQQFKDFQRRILVATNL
				†		FGRGMDIERVNIVFNYDMPEDS
						DTYLHRVARAGRFGTKGLAITF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22299	52667	Α	22427	9	1535	EAQQLVSERPAENRSWKCLLA
ľ			·			ARGEERGASIMAEQDVENDLL
						DYDEEEEPQAPQESTPAPP\KKD
						IKGSY\VSIHSSGFRDF/LCLKPE
						LLRAIGGLVAFEHPSEVQHE\CI
						PQGHPGAWDVLLPGQSPGMGK
						TAVFRCWPPLQQI\EPVNGQVT\
						VLVM\CHTRELAFQISKEYERFS
						KYMPSVKVSVFFGGLSIKKD\E
						EVMKKNCPHVVVGTPGRILAL
1						VRNRSFSLKNVKHFVLDECDK
						MLEQLDMRRDVQEIFRLTPHEK
						QCMMFSATLSKDIRPVCRKFM
						QDPMEVFVDDETKLTLHGLQQ
						YYVKLKDSE\KTRKVFDLWD\V
						LEFNQVDILRQGQCKRCMAL\A
						QLL\VEQNFPAQWPI\HRGNGPQ
j						KERPVNAYQASSKDFPSGRDPL
						VATQILFG\RGMDIEAKSNI\VFN
						Y\DMP*GLGTPYL\HRV\ARRGR
						FGTKRP*PFTFVSDG\NDA\KILN
						DV\QDRF*S*MWAEL\PEEIDIST
						YIEQSRLNHQLARAGPPGAAPH
						AASPPLSRRHC
22300	52668	Α	22428	3	290	
22301	52669	Α	22429	1	690	MNQTSEKEIISKEQGNVKAAGA
						SDVVLYKIDVPANRYDLLCLEG
						LVRGLQVFKERNFFLTVLEVEK
į						SKTKVEQIQCLCSQEDIADKLG
Ì						VDISATKAVHISNPKTAEFQVA
						RTTLLPGLLKTIAANRKMPLPL
						KLFEISDIVIKDSNTDVGAKNYR
						HLCAVYYNKNPGFEIIHGLLDRI
						MQLLDVPPGEDKGGYV/IQSVG
						KLGVLHPDVITKFELTMPCSSL
						EINIGPFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22302	52670	A	22430	2	1834	GTRPTVSVKRDLLFQALGRTYT
		1				DEEFDE\LCFEFGLELDEITSEKE
			į			IISKEQGNVKAAGASDVVLYKI
		ļ				DVLATRYDLLCLEGLVRGLQV
-		1				FKERIKAPVYKRVMPDGKIQKL
				1		IITEETAKIRPFAGAAVLRNIKFT
						KD\RYDSFIELQEKLHQNICRKR
						ALVAIGTHDLDTLSGPFTYTAK
						RPSDIKFKPLNKTKEYTACELM
						NIYKTONHLKHYLHIIENKPLYP
						VIYDSNG\VVLSMPPIINGDHSRI
						TVNTRNIFIECTGTDFTKAKIVL
						DIIVTMFSEYCENQFTVEAAEV VFPNGKSHTFPELAYRKEMVR
ļ						
1						ADLINKKVGIRETPENLAKLLT
		l				RMYLKSEVIGDGNQIEIEIPPTR
						ADIIHACDIV\EDAAIAYG\YNNI
			1			\QMHSPENFTP*ANQFPLNK\LT
						ELLRHDMA\AAGFH*STLPLAL
					}	CSQGRYWLDKLGVGYLWQQR
						AVHISN\PKTAEFQVARTTLLPG
]				PP*RP*AANRKV\PLPL\KLFEIS\
						DIVIK\DSNT\DVGAKKTTRQSL
		l				VAGLLPTKNPG\FEINSWGCWD
						RIMAVCSDV\PPGGRQGGDM*F
		1				KASEGPAFFPGAMCRDSFARGQ
						SL\GKLG\VLHPDVIHQIWS*PLP
						CSSLEI\NIGPFL
22303	52671	A	22431	11	304	
22304	52672	A	22432	1	642	
22305	52673	В	22433	1	423	LUCUDATECE ALCOVER OF CLASS
22306	52674	Α	22434	1	525	LHEHPVTESPMNCSVTLQLSLY
		1				MKSVA/GIGQTGKIEPVVLPLL
						WFAESGAMEGETLHTFYTQLV
						LMPKVMHYAQYVLLALGCVL
						LLVPVICQIRSQ/DRCWPEGSPG
				1	1	*QPFAFPPA*TDIILWTEPATLM
						AEVSACCAPELPPPPHRVPQIIG
		1				AQAEVNEPGGPAFRALLDPGA
22307	52675	Α	22435	3	483	VRGSGNSHGEVSKGRMCLAGA
		1		1		LPLPGACAALVLLLVILPEPLRC
		F		1		PLHTLYGGRDPVDSIRLWRQA
		-		1	:	GEFRHKSNITFNNNDTVSFLEY
		1				RTF\QFQPSKSHGSESDYIVMPN
]				ILVLNKPRSTWISVCFVHGCIPA
		1				SRTGPRTESMHSKSLWKAWLA
1	I	1	I		I	APGLPLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22308	52676	A	22436	1	1623	RRRPRLLPGAEPCEPRVGPRRA
						DMGCSAKARWAAGALGVAGL
		ŀ				LCAVLGAVMIVMVPSLIKQQV
		ŀ				LKNVRIDPSSLSFNMWKEIPIPF
						YLSVYFFDVMNPSEILKGEKPQ
		ļ				VRERGPYVYK/ESSRHKSNITFN
						NNDTVSFLEYRTFQFQPSKSHG
1		1				SESDYIVMPNILVLGV\AVMME
		1				NKPMTLKLIMTL\AFTTLG\ERA
		ł				FHGTALWGEV\MWGYKDPLVN
,				i		LINKYFPGMFPFKGGFGVFAEL
						NNSDSGLFTVFTGVQNISRIHLV
						DKWNGLSKVDFWHSDQCNMI
						NGTSGQMWPPFMTPESSLEFYS
		İ				PEACRSMKLMYK\ESGVFEGIP
						TYRFVAPKTLFANGSIYPPNEGF
						CPCLESGIQNVSTCRFSAPLFLS
						HPHFLNADPVLAEAVTGLHPN
						QEAHSLFLDIHPVTIIPMNCSVK
						LQLSLYMKSVAGIGQTGKIEPV
						VLPLLWFAESGAMEGETLHTF\
						YTQLVL\MPKVMH\YAQYVPS
						WALG\CVLLLVPV/LSCQIRSQE
		i				KCYLFWSSSKKGSKDKEAIQA
						GSE*SLMTSAPKGS\VLQEAKL
22309	52677	Α	22437	198	1062	STCVEMPRNLLYSLLSSHLSPHF
						STSVTSAKVAVNGVQLHYQ\HF
						GEGDHAV\LLLPGMLGSGETDF
		1				GPQLKNLNKKLFTVVAWDP\R
						GYGHSRPPDRDFPADFFERDAK
						DAVDLMKALKFKKVSLLGWS
						DGGITALIAAAKYPSYIHKM\VI
					}	WG\ANAYVT\DERPAG*YEGHP
						RMVSQMGVEGKKESL*EALYG
			1			YDYFARTCEKWVDGIR\QFKHL
]		PDGNICRHLLP\RVQ\CPPPLIVH
				1		G\EKDPLVPRFHADFIHKHVKG
						SRLHLMPEGKHNLHLRFADEF
		<u> </u>				NKLA\EDFLQ
22310	52678	Α	22438	223	393	AQASPPLLPP**ERSNVPCPIS

2902

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	P. Company of the Com	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22311	52679	Α	22439	524	1516	FILEDITGYVMAPIDGLLCREPR
						RGLYLEGTKPMTLFQIQFLNSV
						GDLLDLIPSLSPTKNGSLKVFKR
						WDMGHCSALIKVLPGFENILFA
		1				HSSWYTYAAMLRIYKHWDFNII
						DKDTSSSRLSFSSYPGFLESLDD
						FYILSSGLILLQTTNSVFNKTLL
						KQVIPETLLSWQRVRVANMMA
	ŀ					DSGKRWADIFSKYNSGTYNNQ
						YMVLGPEESKAEPQSLTKGT/P
						VTLWEQIPTY/VRNISGTNLMFL
						RGKGFWA\SYNVPFHEKIYNWS
						GLSHC*FRKLGLGPTPYDFRPH
						EPKIFRRDPR/DK*LIRHPMKYI
						MRYN\NYKE\DPYSRGDPPVNT
22312	52680	A	22440	1	306	
22313	52681	Α	22441	698	1613	FQFFQIFPEYKNNDFYVTGESY
						AGKYVPAIAHLIHSLNPVREVKI
						NLNGIAIGDGYSDPESIIGGYAE
						FLYQIGLLDEKQKK\YFQKQCH
			!			ECIEHIRKQNWFEAFEIL\DKLL\
						DGDLTSDPFLTSQNVTGCS\NY
						YNFLRCTEPEDQLYYVKFLSLP
						EVRPAHPTLGN\QTFN\DGPIV*
						KVPCGEDTSTSPVKPW\LTEI\M
						NNYRVLIY\NG\QLGHPSVAAAP
						DQSAPLMG\MDWKGSP\EYKK
						AEKK\VWKIFKS\DSEVA\GYI\R
						QVGDF\HQVIIRGGGHILPYDQP
22214	50.600	ļ	22442	1	250	LRAFDMINRFIYGKGWDPYVG
22314	52682	A	22442	1	258	
22315	52683	A	22443	366	848	ECLOVETA VEIBLIEL VEMLTEL
22316	52684	Α	22444	L	585	ECLCVSTAVRIRHELKTMLTEL LRGGSFEFKDMRVPSALVTLH
						l I
						MLLCSIPLSGRLDSDEQKIQNDI
						IDILLTFTQGVNEKLTISEETLA
		1				NNTWSLMLKEVLSSILKVPEGF FSGLILLSELLPLPLPMQTTQVS
						LPYNMHLINDCSNTF*KASDSV
						KKQKPSSPFLPKRHKITG*GGD
						l `
L	1	1	<u> </u>		<u> </u>	KTETSWSLRKYGGKNFK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22317	52685	A	22445	625	2700	YIRGHSTPRKLWSMHLHVQAK
						LLQEIVRSFSGTTCQPIQHMLRR
						ICVQLCDLASPTALLIMRTVLD
						LIVEDLQSTSEDKEKQYTSQTT
						RLLALLDALASHKACKLAILHL
						INGTIKGDERYAEIFQDLLALVR
						SPGDSVIRQQCVEYVTSILQSLC
						DQDIALILPSSSEGSISELEQLSN
						SLPNKELMTSICDCLLATLANS
İ						ESSYNCLLTCVRTMMFLAEHD
						YGLFHLKSSLRKNSSALHKLLK
						RVVSTFSKDTGELASSFLEFMR
						QILNSDTIG\CCG\DDNG\LMGSR
						RGAHTSRTMSINAAELNQLLQS
						TEESPEDLFLELEKLDLEHSKD
						DDNLYSLLD**RGLKQMLESSG
						DPLPLSDQDVEPVLSAPESLQN
						LFNNRTAYVLADVM\DDQLKS
						MWFTPFQAE\EIGTDLDLVKVD
						LIELSEKCCSDFDLHSE\LERSFL
						SEPSSPGRTKTTKGFKLGKHKH
						ETFITSSGKSEYIEPAKKEFMLC
	:					PPPRGRGRGGFGQGIRPHDIFR\
						QRKQNTSRPP\SMHVDDF\VAA
						E\SKEVVPQDGI/PSHPNRPLKVS
						QRISSRVGFSGNRGGRGAFHSQ
						NRFFTPPASKGNYSRREGTRGS
						SWSAQNTPRGNYNESRGGQSN
1						FNRGPLPQL\RPLKFPTG*PAPK
						SLGTRAS*MVRGG\LGTFPGLS
						AKLAASGGSRGKFVSGG\SGRG
22318	52686	Α	22446	2487	3161	TWWLHISIIIPGGKKSYSPRLGR
1						GKRFATFVLLVGGELCLTSYAP
1						PSIHPILRSEAECLNYSIFSKFLA
		3				PAGEFALPRSFDNCPRVRLRAL
						RLALLILLTFILCWTPYYLLGM
[WYWFSPH\MLTEVPPSLSHILFL
1						LGLLNAPLDPSPSYGAFTLGCR
		Ì				RGHQELSIDSSKEGSGRMLQEEI
	[HAFRQLEVQKTVTSRRAGETK
						GISITSI*S*QSM*EQNSKSLVP
22319	52687	Α	22447	232	337	
22320	52688	Α	22448	1	1449	
22321	52689	Α	22449	1	2004	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22322	52690	A	22450	3	1558	 APARDRAGGGRRALPRLPAAS
22322	32070	1	22 130	ا ا		ARARGLTRALSRAMACRQEPQ
						PQGPPPAAGAVASYDYLVIGGG
						SGGLASARRAAELGARAAVVE
						SHKLGGTCVNVGCVPKKVMW
						NTAVHSEFMHDHADYGFPSCE
						GKFNWRVIKEKRDAYVSRLNA
						IYQNNLTKSHIEIIRGHAFF\TSD
						PKPTIEVSGKKYTAPHILIATGG/
						MVPSTPH*EPRSRGASL*ITSDG
						FFQLEEIVPAAASLVGARLPLSV
						EMAGIL\SALGSKTSLMIRHD\K
						VLRSFDSMISTNCTEELENAGV
						EVLKFSQVKEVKKAL\SGLEAS
						MVTAVPGRLPVMPMIPDVDCL
						LWAIGGVPNTKDLSLNKLGIQT
						DDKGHIIVDEFQNTNVKGIYAV
]			GDVCGKALLTPVAIAAGRKLA
			1			HRLFEYKEDSKLDYNNIPTVVF
						SHPPIGTVGLTEDEAIHKYGIEN
						VKTYSTSFTPMYHAVTKRKTK
						CVMKMVCANKEEKVVGIHMQ
į						GLGCDEMLQGFAVAVKMGAT
						KADFDNTVAIHPTSSEELVTLR
22323	52691	A	22451	51	464	
22324	52692	В	22452	103	1184	
22325	52693	Α	22453	52	148	
22326	52694	Α	22454	1	3282	
22327	52695	Α	22455	145	436	
22328	52696	Α	22456	1	3622	MPGYRAGPFEEAFDREDERQIE
						CQGRGVDRKITGEHTTFSGRSG
						PDHELTLDSGKGCTRSMVPASA
						ASEDRRKLPIIVEDEGGPTRSRA
						CSSPARGSRPPPSAIGCSPVAQA
						SDSAAGPARRTALQSLSSWLGY
					•	QIDRHSVPVYVFKSPLFSVIMAP
						KHKSSDAGNLDRPKRSRKVLPL
						SEKVKVLDLIRKDKKSYAEVA
		1				KIYGKNESSIREIVKKEKEIRASF
				1		AVSPPTAKVTATVRDKCLVKM
		1	<u> </u>			EQALHLWVEE
22329	52697	Α	22457	3	422	VVELRCQDKEGNPAPEYTWFK
						DGIRLLENPRLGSQSTNSSYTM
						NTKTGTLQFNTVSKLDTGEYSC
						EARNSVGY/PQVPGKRMQVDD
						LNISGIIAAVCGLGVCYAQRKG
			1			YFSKETSFQKSNSSSKATTMSE
		 				NDFKHTKSFII
22330	52698	A	22458	1	208	
22331	52699	A	22459	122	321	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22332	52700	A	22460	2	1169	HTPKTEQTPIPGLEDPPLGSQLR
22332	32700	^	22400	2	1109	RRPGEGETDIPSRAVPPLPPLPT
						ł
						LCSFPAPEVQGPPASCAPAAGT
						LDLLRAAGCRPGKMARRSRHR
						LLLLLRYLVVALGYHKAYGF
						SAPKDQQVVTAVEYQEAILAC
						KTPKKTVSSRLEWKKLGRSVSF
İ						VYYQQTLQGDFKNRAEMIDFNI
						RIKNVTRSDAGKYRCEVSAPSE
						QGQKPG\EDTVTLEVLGDVHVL
						APAVPSCEVPSSALSGTVVELR
İ						CQDKEGNPAPEYTWFKDGIRLL
						ENPRLGSQSTNSSYTMNTKTGT
						LQFNTVSKLDTGEYSCEA\RNS
						VG*SEGPCETNASR*SQHKVGII
						GSRS*LWALVDFPFVGLVYSYG
						SEGKATFSKETSFQKSNSSSKAT
00000	50501	<u></u>	20.461	67	1/0	TMSENDFKHTKSFII
22333	52701	В	22461	57	168	
22334	52702	A	22462	1 200	682	CTEDDSCLITCKKDKALLCACD
22335	52703	A	22463	398	1465	GTFPPSGLITGKKRKALLGACD
					1	CTQIVKPSGVHLKLVLRFSDFG
				Ì	ļ	KAMFKPMRQQRDEETPVDFFY
						FIDFQRHNAEIAAFHLDRILDFR
						RVPPTVGRIVNVTKEILEVTKN
		1				EILQSVFFVSPASNVCFFAKCPY
						MCKTEYAVCGKPHLLEGSLSA
		1				FLPSLNLAPRLSVPNPWIRSYTL
						AGKEEWEVNPLYCDTVKQIYP YNNSQRLLNVIDMAIFDFLIGN
						MDRHHYEMFTKFGDDGFLIHL
						DNA\RGFGRHSH\DEIPILSLLSQ
						CGMIKKNTLLHLQ\LLAQADYR
						LS\DVMRESLLEDQL\SPVLTEP
						HLLAL\DRRLQTILRTV\EGCIV\
						AHG\QQSVH*FDGPSGNSSGPR
22336	52704	A	22464	2	424	GRVGVSAVIVRAWITHKYSGR
2230	32/04	^	22704	-	T	DWNVKWEENLLHAVAKNYTL
		1				LQTIPPFERPFKDHQVCLEWNM
		1				GYIWNLRANRIPQCPLEN/DFRE
		1				DFHYNDTAGYFIIGGSRYVAGI
		1		1		EGFFGPLKYYRLRSLHPAQIFNP
						LLEKQLAEQIS
22337	52705	A	22465	23	323	Provention
22338	52706	A	22466	3	287	LSVQMYAQAALDGDSQGFFNL
	32700	[,		ľ		ALLIEEGT\YERCWSHSNEESFS
		1			•	PCSLAWLYLHLRLLWGAILHSA
		1			1	LIYFLGTFLLSILIAWTVQYFQS
		1			1	VSGKDL
	 	+	22467	1,	240	
22339	52707	IA	22467	11	348	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22341	52709	Α	22469	1	507	
22342	52710	Α	22470	187	3682	SILCLLSPCVVQFGKPVVSILSS
						RSRHSPCTKKGWEGMRKHLHT
	v				·	RQGHK*VHV\KFP\SIMGLQGM
]			DYFHPDISIFCNLAVIVRAW\ITH
			ľ			KYRGRDWNVKWEENLLHAVA
:						K\NYTLLQTIPPFERPFKDHQVC
					·	LEWNMGYIWNLRANRIPQCPL
<u> </u>						ENDVVALLGFPYASSGENTGIV
		İ				KKFPRFRNRELEATRRQRMDYP
					:	VFTVSLWLYLLHYCKANLCGIL
				1	;	YFVDSNEMYGTPSVFLTEEGYL
		l	1			HIQMHLVKGEDLA
22343	52711	Α	22471	3	153	
22344	52712	Α	22472	1	408	TEPEFPGRRFRPRMAVMAP\RT
						LVLLLSGALALTQTWAGSHSM
		1				RYFSTSVSRPGRGEPRFIAVGYV
		1				DDTQFVRFDSDAASQRMEPRA
		-		1		PWIEQEGPEYWDEETGKVKAH
						SQTDRENLRIALRYYNQSEAGS
						HTLFPR
22345	52713	Α	22473	1	823	MAVVAPRTLLLLLSGALALTQT
						WAGSHSMRYFSTSVSRPGSGEP
				,		RFSAVGYVDDTQFVRFDSDAA
•						SQRMEPRAPWMEQEEPEYWDR
						QTEISKTNAQIDLESLRIALRYY
						NQSED/GPPPKTHMTHHPISDHE
						ATLRCWALSFYPAEITLTWQRD
						GEDQTQDTELVETRPAGDGTF
ļ						QKWASVVVPSGQEQRYTCHVQ
						HEGLPKPLTLRWEPSSQPTIPIV
		1			}	GILAGLVLFGAVIAGAVVAAV
						MWRRKSSDRKGGSYSQAASSD
						SAQGSDVSLTACKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22346	52714	A	22474	1	1162	NSRVDDFVAVMAPRTL\LLLLS
						GALALTQTWAGSHSMRD\FFTS
	1					VSRPG\R*EPRFIAVGYVDDTQF
						VRF\DSDAASQRMEPRAPWIEQ
						EGPEYWDQETRNMKA\HSQTD
						RENLRIALPLTYNQSEAGS\HTL
						QMMFGCERGGSDGALSFRGLP
						PVTAY\DGKGFTFALNE\DLRS
						WT\AADMAAQITQRKWETPPM
[1				WRS/EQEEAYLEG\QCV\DGLRR
						YL\ENGKEDAAAQRTPPKTHMT
						HHPIS*P*RPL*RCW\AL\AFYPC
		1				GDHT*PWQRDG\EDQTQ\DTEL
		ŀ				V\ETRPAGDGTF\HKWAAVVVP
						SGEESEITPANVQQ*RVLPKPLT
						LKM/WEPSSPAHPSPIVGHHLLA
						LVL/LLGAFDHLKLWVAAVML
1						KEGTAQVGKG*KIEKGGSYSSR
						LPSQ
22347	52715	A	22475	9	316	IPAMGLETEKADVQLFMDDDS
		-				YSHHSGLEYADPEKFADSDQD
						RDPHRLNSHLK\ILMPFVKTCL
						MVLPSVQTIWKSVTDVIIAPLCT
						SVGRCFSSVSLQLSQD
22348	52716	A	22476	3	606	RPRTKAAMGLETEKADVQLFM
						DDDSYSHHSGLEYADPEKFADS
						DQDRDPHRLNSHLKLGFE\DVI
						AEPVTT\HSFDKVW\ICSHALFEI
						\SKYRM\YRSLT\VFLGHSPWAFI
						CGEFSFATL\SCLAHLGF*WPFV
						\KTCLM/VFLPSVQTIWEEC*QM
	+			1		V/ISLAPLC\TSVG\RCFLFLFSLA
						TEPRNGILGPQVWRLGYCNTSF
				1		VIIT
22349	52717	A	22477	101	432	ATRDQLKAKPENLLGSISRDSG
				1		RTQEGIPARVPYPPPP/PTREPPQ
						ST*PGRNGLLGCLGGGAPRPG
ļ						WLDSEAGGAPAFHCPG*CPRRH
ĺ						RRGGTRLKQLPPPPTTVHDQPL
22350	52718	A	22478	794	1209	VKVDEVTVKHLGCQRCQWSRC
				1		PPSCSGTCLCCGAEAPCAAQPA
						WPPPPAAPPPTPPAPPSD\NSSCT
						Q*PTPRCGT*DKDKGIFTLPKGH
						P*V*SLEHGFLKLGSCP\PPPPHA
						LVRPPQVRFNTPHGAEPQRDQL
						1
						1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	họđ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ				sequence		
22351	52719	A	22479	3	1041	GSNCPLRQADPGPPRHTLFMDF
		-				VAGAIGGVCGVAVGYPLDTVK
						VRIQTEPKYTGIWHCVRDTYHR
						ERVWGFYRGLSLPVCTVSLVSS
					}	VSFGTYRHCLAHICRLRYGNPD
		l]	AKPTKADITLLGCGLRTRPRVP
						DVAH*GGQSPLADADTGFLSSS
						GGFRLSALLAAPPMCPVPPACP
						EPKYRGPLHCLATVAQ\EEGLC
						GLYKGSLALVLRDGHSFATYFL
1						SYAVLCEWLSPAGHFRPDVPG
						VLVAGGCAGVLAWAVATPM\D
		ĺ				GIKSRLEANGQGQ\NGYRGLLH
		l				CMVTSVREEGPRVLFKGLVLN
•				}		CCRAFPVNMVVFVAYEAVLRL
		ŀ				RPGSAHHSRSPTPRGATHQQLV
22352	52720	A	22480	38	453	RFG3AHH3K3F1FKGA1HQQLV
22353	52721	A	22481	287	413	DMDYRRGGWTVIHRKIGGRID
22333		` `		20,		FQRVR*DYLHGYADLLREFWL
22354	52722	A	22482	1	393	MPKLLKWNYEGGACWGNLW
						KLPNVERLCFSEATVFRSAAQC
]		1				LVVGLGLLLVSRACSHEDAIEW
		1				RGDSSVVNIVEDGSNAKDESKS
						NDTVCKEDCEESCDVKTKITRE
		l				EKHFMCIFHNVQSLGTVCGEK
						GIDVN*NYTRRKTFHVYLPQCS
						VTGNSLRGERH
22355	52723	A	22483	505	854	
22356	52724	Α	22484	584	1009	FTQKDLFIPFEVMCDMDYRGG
1						GWTVIQKRIDGIIDFQRLWCDY
						LDGFGDLLGDAFL\FL*KEDNQ
į						NAMPFSTSDVDNDGCRPACLV
					,	NGQSVKSCSHLHNKTGWWFN
[KCGLANLNGIHHFSGKLLATGI
						QWGTGNQKQLTWQE
22357	52725	Α	22485	1	3123	MSHRARPCPAPFLFWMLGVKT
		ļ				EPAGQDRRKKARSRCCIFRVHA
				1		SGLWGLRFPSVEWGKPSRDKA
						QLSESPPQSVYLFAETALSAIVP
						GDGVAGGGGTGDGGPGAPYPG
						AVADMEELTIWEQHTATLSKD
						PRRGFGIAISGGRDRPGGSMVV
						SDVVPGGPAEGRLQTGDHIVM
	}					VNGVSMENATSAFAIQILKTCT
						KMANITVKRPRRIHLPATKASP
						SSPGRQDSDEDDGPQRVEEVD
						QGRGYDGDSSSGSGRSWD
22358	52726	A	22486	191	267	
	•			·		J

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22359	52727	A	22487	2	609	PPPRSSEDPAKMAEVEHEEGSG
						PFRKFTLPAAVDL\DQ\RLDISY
		[DQLEQPMQAVPVARPAAGGLN
		1				PGACGRKQ\HSPA*KRLLQGPS
						KGGAAPCHKPEAGEDATLRD
	İ	1				MII\LPEMWGS\MVGVYH\GKTF
						\NQVEDQGVCGRPCRLGWAGV
						A*CRRDQLTPSFALGLPQPEMI
· ·		ì				GHYLGEFSITYKPVKHGRPGIG
						ATHSSRFIPLK
22360	52728	A	22488	3	461	STISLYCGLKTCPKLTCAFPVSV
						PDSCCRVCRGDGELSWEHSDG
						DFFRQPATREASHSCHRSHYDP
						PPSRQAGGLSRFPGARSHRGAL
						MDSQQASGTIVRFVLNNKHKH
						GQVCVSNGRTYSHGESWHPNL
						RAFGIVECVLCTPTPID\WPQGD
						*GKCRHGCPTLHYVD*LPLEKQ
						VILATALTMILHQADRLEVCPA
						FLGPEVTGELLWIPSKHQEPLC
						DLSSITNTSMDKCVFPMEGPIL
						MASPGTQTSGLLALWSVCYVL
						PPPSIVPQGD
22361	52729	A	22489	656	1495	TGDGELSWEHSDGDIFRQPANR
			ļ			EARHSYHRSHYDPPPSRQAGGL
						SRFPGARSHRGALMDSQQASG
						TIVQIVINNKHKHGQVCVSNGK
						TYSHGESWHPNLRAFGIVECVL
						CTCNVTKQECKKIHCPNRYPCK
						YPQKIDGKC*QNSTCFLIPEELP
						GQSFDNKGYFCGEETMPVYES
		İ				VFMEDGETTRKIALETERPPQVI
		1				CSTFCLA\QGILQHFHIEKISKR
		ļ				MFEELPHGHFAFVCLLGQWKIF
		l				TEGEAQISQMCSSRVCRTELED
						LVKVLYLERSEKGHC
22362	52730	Α	22490	1	477	
22363	52731	Α	22491	1	294	
22364	52732	Α	22492	56	4042	VFFTSMGEHLQKLIPASPACLA*
		ł				IPQACQP*PPTPPAWQCRNLR/L
						AASRHQEPRGEVGGPLGHGAE
						RILAVPVRTDAQGRLVSHVVSA
				1		ATSRAGVRARRAAPVRTPSFPG
		1				GNEEEPGSHLFYNVTVFGRDLH
				1		LRLRPNARLVAPGATMEWQGE
		1		1		KGTTRVEPLLGSCLYVGDVAG
						LAEASSVALSNCDGLAGLIRME
						EEEFFIEPLEKGLAAQEAEQGR
						VHVVYRRPPTSPPLGGPQALDT
		1				GASLDSLDSLSRAL

	of peptide				1	Amino acid sequence (X=Unknown,
i 16		hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22365	52733	Α	22493	173	1148	QALGLPSLLAVLALSTTDRASF
		į .				SAVISRSPQRTVSSTRKEQRPLV
						FAFSALRLGTNACKMDGFCSTF
				1		LGAPGIRLKLLTPAGTHKSERD
						RTATPTRHHEKRLFCSSLSIGRQ
						AEEAWRADHPLPESRLFGLVPL
						GGAAWCRLAGSFCVCSLRLPRP
						IELVRAASGALLGAVRRRNHGP
						PHCFSTATLLCKAYRGGHLTIR
						LALG\GAPNRPFYRIVAAQNKC
						PRDGPFP*SKLGPPYDP/LLPPNS
						HGEKTPCPLN\LDRI\RHWIG\CG
		İ		!		APPLLSPMEKLLGLAGFFP\LHP\
						MMITNAERLRRKRAREVLLAS
						QKTDAEATDTEATET
22366	52734	A	22494	2	4985	AGATTALELPGQPSVTGVPELP
						GLPSATRALELSGQPVATGALE
						LPGPLMAAGALEFSGQSGAAG
						ALELLGQPLATGVLELPGQPGA
						PELPGQPVATVALEISVQSVVT
					İ	TSELSTMTVSQSLEVPSTTALES
						YN/DERSMMSSYERSMMSYER
						SMMSPMAERSMMSAYERSMM
						SAYERSMMSPMAERSMMSAYE
						RSMMSAYERSMMSPMADRSM
						MSMGADRSMMSSYSAADRSM
						MSSYSAADRSMMSSYTADRSM
22367	52735	A	22495	18	174	ALIKKHFIF*IAAAKPTPPKSQV
						TLTKEFPVSSGSQHRKKEADSV
						YGEWVL
22368	52736	Α	22496	3	595	IKKHFIF*IAAAKPTPPKSQVTLT
						KEFPVS\SGPQHRKKEADSVYG
						EWVPVEKNGEENKDDDNVFSS
						NLPSEVSRRNIMYFSFFIPESAH
						FIVMLCLIWDSVSLFLRAGVKR
						QGRV*DGQM\KQPAASH\LTVT
						RM/CNSLCG\TKATKWKRLRFA
						ENSVI\TSLPNIGPSLHLWEGSP
						WYSIPFSFPSLLLLKVGEVEFKR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
22369	52737	A	22497	3	5618	SSELPGPSVTPVPQLSQELPGLP
22307	32/3/	``	2277	٦	3070	APSMGLEPPQEVPEPPVMAQEL
						PGLPLVTAAVELPEQPAVTVA
						MELTEQPVTTTELEQPVGMTTV
						EHPGHPEVTTATGLLGQPEATM
						VLELPGQPVATTALELPGQPSV
]				TGVPELPGLPSATRALELSGQP
						VATGALELPGPLMAAGALEFS
						GQSGAAGALELLGQPLATGVL
						ELPGQPGAPELPGQPVATVALE
						ISVQSVVTTSELSTMTVSQSLEV
						PSTTALESYNTVAQ
22370	52738	A	22498	1	4685	MAPESHILASNTMETHILASNT
				-		MDSQMLASNTMDSQMLASNT
						MDSQMLASSTMDSQMLATSSM
						DSQMLASNTMDSQMLASNTM
						DSQMLASSTMDSQMLATSSMD
						SQMLATSSMDSQMLATSTMDS
						QMLATSSMDSQMLATSSMDSQ
						MLATSSMDSQMLATSSMDSQM
						LATSTMDSQMLATSTMDSQML
						ATSSMDSQMLASGTMDSQMLA
						SGTMDAQMLASGTMDAQMLA
						SSTQDSAMLGSKSPDPYRLAQD
						PYRL
22371	52739	Α	22499	291	7655	KRAPEKVDAYLYKQILLMKLP
						TKKSKKHKKHKNKKKKKKKE
						KEKKYKRQPEESESKTKSHDD
						GNIDLESDSFLKFDSEPSAVALE
		İ				LPTRAFGPSETNESPAVVLEPPV
			1			VSMEVSEPHILETLKPATKTAE
						LSVVSTSVISEQSEQSVAVMPEP
						SMTKILDSFAAAPVPTTTLVLK
						SSEPVVTMSVEYQMKSVLKSV
						ESTSPEPSKIMLVEPPVAKVLEP
						SETLVVSSETPTEVYPEPSTSTT
			1			MDFPESSAIEALR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22372	52740	A	22500	90	2166	SQHFGRLRRVDYLRTSKMQTSS
						SRSVHLSEWQKNYFAITSGICT
						GPKADAYRAQILRIQYAWANS
						EISQVCATKLFKKYAEKYSAIID
						SDNVESGLNNYAENILTLAGSQ
						QTDSDKWQSGLSINNVFKMSS
						VQKMMQAGKKFKDSLLEPALA
						SVVIHKEATVFDLPKFSVCGSS
		1				QESDSLPNSAHDRDRTQDFPES
						NRLKLLQNAQPPMVTNTARTC
						PTFSAPVGESATAKFHVTPLFG
						NVKKENHSSAKENIGLNVFLSN
						QSCFPAACENPQRKSFYGSGTI
						DALSNPILNKACSKTEDNGPKE
						DSSLPTFKTAKEQLWVDQQKK
						YHQPQRASGSSYGGVKKSLGA
ļ	•					SRSRGILGKFVPPIPKQDGGEQN
	:					GGMQCKPYGAGPTEPAHPVDE
						RLKNLEPKMIELIMNEIMDHGP
		l				PVNWEDIAGVEFAKATIKEIVV
						WPMLRPDIFTGLRGPPKGILLFG
		ļ				PPGTGKTLIGKCIASQSGATFFSI
						SASSLTSKWVGEGEKMVRALF
						AVARCQQPAVIFIDEIDSLLSQR
						GDGEHESSRRIKTEFLVQLDGA
1						TTSSEDRILVVGATNRPQEIDEA
1						ARRRLVKRLYIPLPEASARKQI
	1				:	VINLMSKEQCCLSEEEIEQIVQQ
						SDAFSGADMTQLCREASLGPIS
						AV/SQTADIATITPDQVRPIAYID
						FENAFITVRPIVSPKDL*LYENW
22373	52741	Α	22501	3	523	FFFFFPAFKMSKRG\RGWVPSG
						CRKFPDFPLGLP\VGTL*SNWCL
						TNT\GAQKTLYI\ISREGGIKGTG
						LTRLPRLPGVG\DMVMAHSSRK
						ANPELQKKGYIPA\VVIRQRKVI
						TVGKRWRCFFIFED\NAGVIGD\
						NKG\EMKGFAITG\PRAKECAD
L						LWPRDCIQMLGSIAWFSQYIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22374	52742	Α	22502	3	946	GPSASVSFWLWSQQHSPRIRHP
						RPGERRRLLHGIPHPTAEKQPA
		ļ				PLRPERRGGRRRRPAGRRGCA*
						/GGARPCSSRSS*PASTPAPRAA
						PPCRSPCSARPRTGPRTGAGRPR
						AP/HDGPPCATPDPRDQAS\PLPP
						RGPRPRVAPGSP\PEGHPRELPS
		ŀ				EGSPGPRAPGQRHVRPPRAVLR
		ŀ				PRGLRGGGCPGPAHPAPRLQRR
						RAPGEAAAPGPRWPGRPGRGR
		ŀ				AQALVPAARSRCFWKLWEAVF
						CLPVAAAPTRRFGARQRPQLW
:						LHGHKGGQVGEVSSHSAGEEH
ı				ŀ		LVPHAAPKRAREPHGRSGSPGG
						IASSWQDGS
22375	52743	Α	22503	34	823	
22376	52744	Α	22504	424	2339	LLRGTLQSYAWTSVIPDIATELR
						DAPEKTLACMGLAIHQVTICLM
						HTFV*TFLMFAHTCTPCLTMVN
		İ				VPHIHAR*GI*CIKVLLRMGH*R
						PFKNENAHRGTVVRVSNIKPLC
						TKMAFLCAACGEIQSFPLPDGK
		l				YSLPTKVIRSLTASLFILVKKAN
						QNTENISRQSIK*AIRLLHKRQA
						LDVTLLRKSIILLHDLVDSCVPG
						DTVTITGIVKVSNAEEGRVQLF SLFVSQ**FINVHLF*DAPPHILV
						VGDPGLGKSQMLQVDNQSFSV
		l				CSFACKMR**FTSEFSPLSKDSS
						SGDFALEAGALVLGDQGERPK
						GIISDSGTF*KAFSFSVNFLSISL
		İ				AKAGVVCSLPARTSIIAAANPV
						GGHYNKAKTVSENLK*VSSNIY
		1				TFNVLNHFNSIRKRNSSHLLSEH
						VIAIRAGKQRTISSATVARMNS
						QDSNTSVLEVVSEKPLSERLKV
						*MFLLLIQFGSV*MSKFSVWLIV
	1					DVYPRLSTEAARVLQDFYLELR
						KQSQRLNSSPITTRQLESLIRLTE
			ŀ			VCFFLWSCFFWLKGEGCALTSL
						LKYR*DKNFKLF*MKYQLSSHL
			}			KLGNRPNRSTAKRFISALNNVA
						ERTYNNIFQFHQLRQIAKELNIQ
					1	VADFENFIGSLND\QGYLLKKG
						PKVY\QLQTM
22377	52745	Α	22505	73	214	
22378	52746	Α	22506	1	503	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22379	52747	<u> </u>	22507	3	1000	ESKLEGKAAQDIKDEEPGDLGR
22319	32/4/	Α	22307	3	1000	1
				1		PKPECEGYDPNALYCICRQPHN
			ł			NRFMICCDRCEEWFHGDCVGIS
			•			EARGRLLERNGEDYICPNCTIL
						QVQDETHSETADQQEAKWRPG
						DADGTDCTSIGTIEQKSSEDQGI
						KGRIEKAANPSGKKKLKIFQPG
		ŀ				PGPVPTQLPVLWQVLEIAVSRSI
1		ŀ				SAFTLLHCISCKVIEAPGASKCI
						GPG\CCHVAQP\DSVYC\SNDCIL
		ŀ				KHAA\ATMKFLSSGKEQKPKPK
1		ŀ			ŀ	EKMKMKPEKPSLPKCGAQAGI
1		ŀ				KISSVHKRPAPEKKETTVKKAV
						VVPARSEALGKEAACESSTP\S
						WAS\DHNYNA\VKPEKTAAPSP
1					1	VTVV
22380	52748	Α	22508	1	2330	MAAAEGKERAGGRASGRGRSE
						PPRPGARASDAGPGGLREVEGT
1						GEHSVVAAGTVPRVGVGAAAR
		ļ				GEERRAGLGECGPGLGGCVAG
						LISGKSVQAFSVGAGRWPLLRV
		ļ				FGGNVTFCSVVSWSLCAYTLPL
						GMDDKGDPSNEEAPKAIKPTSK
						EFRKTWGFRRTTIAKREGAGD
				ĺ		AEADPLEPPPPQQQLGLSLRRS
						GRQPKRTERVEQFLTIARRRGR
1		ł				RSMPVSLEDSGEPTSCPATDAE
		l				TASEGSVESASETRSGPQSASTA
1		1				VKERPASSEKVKGGDDHDDTS
		1				DSDSDGLTLKELQNRLRRKREQ
						EPTERPLKGIQSRLRKKRREEGP
						AETVGSEASDTVEGVLPSKQEP
	Ì					ENDQGVVSQAGKDDRESKLEG
1				1		KAAQDIKDEEPGD\LGRPKPEC
						EGYDPNALYCICRQPHNNRFMI
				}		CCDRCEEWFHGDCVGISEARG
	·					RLLERNGEDYICPNCTILQVQD
					,	ETHSETADQQEA*MGDLGDA\D
				1		GTDF\TSIGNIRSRSLSRRPKGLK
						GRIEKAANPSGKKELKIF\QPVL
						EIAVSRSISAFTLLHCISCKVIEA
				1		PGASKCIGPGCCHVAQPDSVYC
						SNDCILKHAAATMKFLSSGKEQ
						KPKPKEKMKMKPEKPSLPKCG
						.
						AQPVQWLLLLSSWQLGTSSGL
						QVTLCQPPFTEHKWLKFSVLHG
						ETAGPPRAHRIRKLNKYWGAC
L						HHCMPLRPLPCLPGWPRSSPSL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22381	52749	A	22509	149	2001	IQPPDPRNMTLAAYKEKMKELP LVSLFCSCFLADPLNKSSYKYE\ AGVGDSVGGRMKASGKTVLT GRERRAQADTV\DLNWCVISD MEVIELNKCTSGQSFEVILKPPS FDGVPEFNASLPRRDPSLEEIQ KKLEAAEERRKRGPDLVIYEIH SPASLPFHSSDASVLGDIPPTTY SNISNSNQELLLMGGSPAPEQP HYEANILERSETTSYDNGSDAA QSFKVDGRCVFYINIPIPSNEGK VLWKAPEIATAKEPTWSPLGST GPRPCQILVVKSPVVHGKGPVL SGCGLDLLEQPTFALCFLVAGA VTITFLSCSGEEAREESAPKAMS ERQARRGGAEKQGAEGRGLQA VTTPELPYRMGLGAAFLLKERS GVLPGSFICEDFTAALKQDEEIG
						TMKGKIREAYFKSFVPTPVAFS GGVMLPTGGTLEMCGNNFGGH KEQAGTVLPAFAIALRVTPSAS LVLRPLNLDQAMLQGSLGLQL ADGLFWDLASIMDWTASRIQPS VSSSMKICQAFPGRPLAILLAQC AYVYGTWMKLEAIILTKLTQE QKTKHHLFSLPCCMPADISYRQ RCLGEPPHCTVWNRHLESLVM MSNSTTSPCSELLIFEVPPVAPE
22382	52750	A	22510	373	990	IQPPDPRNMTLAAYKEKMKELP LVSLFCSCFLADPLNKSSYKYE ADTVDLNWCVIFRHWKVIE\LN KCT/SGANSFEVILKPPSFDGVP EFNASLPRRARPIPWEEIPVRKL GRRLRERREVTREADP*KQLAE KR\EHEERGDSKRAFRETTTFIK MAK/EKNLAQKMESNRENREA HLAAMLERLQEK\DKHAEEVR KNKELKEEASR
22383	52751	A	22511	524	1107	LCSPKVLGLNASCQKNFLKQK KSIKTLSTSQ/YSVSGYSLARSE NAHPEFRMAMKDGELVIWDSV HPCYTVFHEQTETFSSLWSEYH DDFRQFLHIYSQDVACYGENLA YFPKGFIENMFFVSANPWVSFT SFDLNVANMDNFFAPVFTMGK YYTQGDKVLMPLAIQVHHAVC DGFHVGRMLNELQQYCDEWQ
22384	52752	Α	22512	1	1413	
22385	52753	В	22513	1	2787	
22386	52754	Α	22514	1	1638	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22387	52755	A	22515	71	2421	YTVCYFLQAAEEDETQNQHDD
22301	32733	^	22313	/ 1	2421	SSSDSGTPSGPDFNYILNMSLW
		l				1
						SLTKEKVEELIKQRDAKGRE\V
						NDLKRK\SPSDLW\KEDLAGIC
						WRDWDKV/ENSPRNEEDVLAG
						MSGKSNLKGKVGKP\KVKKLQ
						LEETMPSPYGRRIIPEITAMKAD
						ASKKLLKKK/KGVDLDTAAVK\
						VEFDEEFSGAP\VEGAGEEGIDS
						HQFPINK\GPKP\RREKKEPGS\R
						VRKTPTSSGKPSAKKVKKRNP
						WSDDESKSESDLEETEPVVIPR
						DSLLRRAADIHRLGGEGFDAGL
					:	TVFAPLFASPFPPRGAPYLKAA
						CQTPRSGIGATAISGEALKGAV
						SSGVDPRWASHLRVKLLEKSFS
						LQEIQSTEKVLSQQAKTLSLLG
		ŀ				EPKLSKQLTGTCIYRKTNLALC
						RGPAAAKSLFQGAVRNDRNKK
						KKETSKQECTESYEMTAELDDL
						TEKIRKAHQETFPSLCQLGKYT
						TSLQKGMQRCQILDLSSVRPIV
						DVRPPELQDSKLVLVQAAKCA
						PLADHCSKPVLLKEGPEDQKD
						QHDVGDCWNPECQTLFDQNN
		İ				AAKKEESETANKNDSSKKLSVE
						RVYHIKTQLEHILLRPDTYIGSV
]	EPLSKVAEEGPAAVGGWKGWF
						GTESRKDPKPTGTEEVERGVW
						LSGREGRLLSPSVVTTMVLAEN
						SAIRFVGGKPSSDTVPKPKRAP
						KQKKVVEAVNSDSDSEFGIPKK
22388	52756	A	22516	1424	4860	GRDRYGVFPLRGKILNVREASH
22300	32730	1	22310	11727	14000	KQIMENAEINNIIKIVGLQYKKS
:						YDDAESLKTLRYGKIMIMTDQ
						DQDGSHIKGLLINFIHHNWPSLL
						KHGFLEEFITPIVKASKNKQELS
						FYSIPEFDEWKKHIENQKAWKI
						KYYKGLG\TSTAKEAKEYFA\D
						MERHRILFRYAGPEDDAAITLA
						FSKKKIDDRKEWLTNFMEDRR
						QRRLHGLPEQFLYGTATKHLTY
						NDFINKELILFSNSDNERSIPSLV
						DGFKPGQRKV
22389	52757	_	22517	87	292	DOLVLOÁVKA
22389	52757 52758	A A	22518	1	903	
	52759	A	22519	3	311	
22391	124127	<u>''</u>	44317	ــــــــــــــــــــــــــــــــــــــ	711	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22392	52760	Α	22520	1	1087	 MPSAAWKRPVEGPATLRRPLGI
		ļ				AAASQCPSSPQPGCLIGAVNLK
		1				SSNRTPVVQEFESVELSCIITDS
		ŀ				QTSDPRIEWKKIQDEQTTYVFF
		1				DNKIQGDLAGRAEILGKTSLKI
						WNVTRRDSALYRCEVVARNDR
						KEIDEIAIEVT\VQVKPVTPVCR
						VPKAVPVGKMATLHCQGE*GP
		ļ				PPA\HYSWYRNDVPLPTDSRAN
		ŀ				PRFRNSSFHLNSETGTLVFTAV
ŀ		ļ				HKDDSGQYYCIASNDAGSARC
ŀ						EEQEMEVCHTDKRGGFPWSW
						AAPHLWFLQDTALLPAALTGW
						CSVSVAFPGAQCKLSVDLSFWG
						LEDGGPLLTAPLGSTPVGTLCR
						GSDPTFPFCTALADDLYESPTPA
						ANFCLDIQAFSYIL
22393	52761	Α	22521	628	1117	FFISVINGQVSSVQRLSGVGPAC
}						LSCGSANPGPPPGTSPGAGAQR
						R*\PRADGSGSPQWPRGARVGG
			-			GRLGTGGRGRPGWRQVPRRLS
			Į.			PGFGR*GGTGPGPVGTSGKRGP
						SRRRAPANDKAACWPRFPGQP
						AS*TGFRGERGVKGFSSWGSG
		ļ				WRAWEDGGTVH
22394	52762	A	22522	2	238	
22395	52763	Α	22523	131	2891	LIGHT LEOD ALONG CONT.
22396	52764	Α	22524	131	3103	LVSFLLFQDALPEGDASPLGPY
						LLPSGAPERGSPGKEHPEERVV
						TAPPSSSQSAEVLGELVLDGTA
		İ				PSAHHDIPALSPLLPEEARPKHA
						LPPKKKLPSLKQVNSARKQLRP
						KATSAATVQRAGSQPASQGLD LLSSSTEKPGPPGDPDPIVASEE
				1		ASEVPLWLDRKESAVPTTPAPL
						QISPFTSQPYVAHTLPQRPEPGE
						PGPDMAQEAPQEDTSPMALMD KGENELTGSASEESQETTTSTIIT
						TTVITTEOAP
	I	<u> </u>	<u> </u>		<u> </u>	IT VITTE QAF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u></u>	<u> </u>				
22397	52765	Α	22525	474	2527	SASPLGPYLLPSGAPERGSPGKE
						HPEERVVTAPPSSSQSAEVLGE
:						LVLDGTAPSAHHDIPALSPLLPE
		1				EARPKHALPPKKKLPSLKQVNS
ļ						ARKQLRPKATSAATVQRAGSQ
					1	PASQGLDLLSSSTEKPGPPGDPD
						PIVASEEASEVPLWLDRKESAV
Ì	1					PTTPAPLQISPFTSQPYVAHTLP
		İ				QRPEPGEPGPDMAQEAPQEDTS
ł						PMALMDKGENELTGSASEESQ
	1					ETTTSTIITTTVITTEQAPGMQPP
						TPMGEGPPIVKHTWILDKCNVA
1						TLQST*ALGPNEGYIDSSDYPLL
						PLNNFLECTYNVTVYTGYGVE
						LQVTPGEYPDGELLSIRGVDGP
) `
						TLTVLANQTLLVEGQVIRSPTN
						TISVYFRTFQDDGLGTFQLHYQ
						GRVRPRLFPRRPDSGDVTVMDL
						HSGGVAHFHCHLGYELQGAK
						MLTCINASKPHWSSQEPICSGM
				ļ		LQPQPHNATIGRVLSPSYPENT
				ŀ		NGSQFCIWTIEAPEGQKLHLHF
						ERLLHDKDR*SSEGPQTNKSA
						LLYDSLQTESVPFEGLLSEGNTI
		1				RIEFTSDQARAASTFNIRFEGEG
				1		PWELYEPYIQNGNFTTSDPTYNI
						GTIVEFTCDPGHSLEQGPAIIECI
				1		NVRDPYWNDTEPLCRGERIHN
						ASAVAGVVLSPNWPEPYVEGE
	1					DCIWKIHVGEEKRIFLDIQ/FVS
						V*NGSVFSQEKVRPTPEWW
22398	52766	A	22526	131	3125	LVSFLLFQDALPEGDASPLGPY
						LLPSGAPERGSPGKEHPEERVV
		ļ			ł	TAPPSSSQSAEVLGELVLDGTA
}						PSAHHDIPALSPLLPEEARPKHA
						LPPKKKLPSLKQVNSARKQLRP
İ		1				KATSAATVQRAGSQPASQGLD
						LLSSSTEKPGPPGDPDPIVASEE
		1				ASEVPLWLDRKESAVPTTPAPL
		1				QISPFTSQPYVAHTLPQRPEPGE
		ı		ŀ		1 -
					1	PGPDMAQEAPQEDTSPMALMD
	•	1				KGENELTGSASEESQETTTSTIIT
	<u> </u>	 				TTVITTEQAP
22399	52767	C	22527	161	352	
22400	52768	Α	22528	2	368	LGSTHASVRINILSVDVLYFIFY
						RRDDNKDIDSEKEAAMEAEIKA
						ARERAIVPLEARMKQFKDMLL
						ERGVRKQSLGEIFLFCISNINMS
						C*KVKSKVSAFSTWEKELHKIV
						FDPRYLLLNLI
		ــــــــــــــــــــــــــــــــــــــ		 	1	

1 -	SEQ ID NO:	,	SEQ ID NO:	1	18	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
22401	52769	A	22529	2	614	KDHRWESGSLLEREEKEKLFNE
		1				HIEALTKKKREHFRQLLDETSAI
		1				TLTSTWKEVKKIIKEDPRCIKFS
		1				SSDRKKQREFEEYIRDKYITAK
						ADFRTLLKETKFITYS*FSERCE
						SESL*KMLGLAAPQTEVLCSST
		1	 			E\RSKKLIQESDQHLKDVEKILQ
	:					NDKRYLVLDCVPEERRKLIVAY
						VDDLDRRGPPPPPTASEPTRRST
22402	52770	В	22530	201	296	
22403	52771	В	22531	1	1290	
22404	52772	Α	22532	3	3369	SSSVMAERGGVGGESERFNPGE
		1				\LRMAQQQALRFRGPAPPP\NA
		İ				AMRGPPPLMRPPPPFGMMRGPP
						PPPRPPFGRPPFDPNMPPMPPPG
		ŀ				GIPPPMGPPHLQRPPFMPPPMSS MPPPPGMMFPPGMPPVTAPGTP
		ŀ				ALPPTEEIWVENKTPDGKVYYY
						NARTRESAWTKPDGVKVIQQS
						ELTPMLAAQAQVQAQAQAQA
						QAQAQAQAQAQAQAQAQAQAQA
						QAQAQAQAQAQAQAQAQAQAQA
						QAQAQAQAQAQAQAQAQAQAQA
						QAQAQAQAQVQAQVQA
22405	52773	Α	22533	43	347	
22406	52774	Α	22534	1	1776	
22407	52775	A	22535	3	433	PLAPNHLGGGAVVLGAESHAS
						KDVAIDMMDSRTSQQLQLIDE
						QDSYIQSRADTMQNIESTIVELG
						SIFQQLAHMVKEQEETIQR*DR
						VNSAHPWDPVHLLPSTQQVFIE
	<u> </u>	↓				HLFM
22408	52776	Α	22536	304	1403	SRGNATGLRTRIRVSTWVSQRH
						RSCPLQLLAVAAATSSPLPPPVT
						LVPPPPDTMSCRDRTQEFLSAC
						KFAARPVKNGI\QTN*GQLLRW
						LVRQRSEIHPSLAKRHLGKDLS
						NTFAKLEKLTILAKR\KPLFDDK
						AGEIEELTYIIKQDINSLIKQIAQ
						LQ\DFVRAKG\SQSGRHLQTHS
						NTIV\VSLQSKL\ASMSNDFKSV
	1					L\EVRDRGTLKQ\QREPEERQFS
						RAPV\SALPLSPTHRAGGA\VVL
					1	GAESHASKDV\AIDMM\DSRTS
						QQLQLIDEQDSYIQSRADTMQN
						IESTIV\ELGSIFQQLAHMVKEQ/
						ERETIQRIDENVLGAQLDVE\AA
						HSEILKYFQSVTSNRWLMDKIF
22.422	50777	ļ	100505	1.42	706	LILIVFFIIFVVFLA
22409	52777	В	22537	147	506	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ļ.	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22410	52778	A	22538	2	1397	AAPGNGLSPSSWRSELACEFQH
						ECTAEVRFGAVVCRGARSHSFY
						LEREILQ/PPGMSL/PENPVTPML
				1		RHLMYKIKSTGPITVAEYMKEV
				1		LTNPAKGYYVYRDMLGEKGDF
						ITSPEISQIFGELLGIWFISEWMA
						TGKSTAFQLVELGPGRGTLVGD
						ILRVFTQLGSVLKNCDISVHLVE
				i.		VSQKLSEIQALTLTKEKVPLER
						NAGSPVYMKGVTKSGIPISW*S
						RICTD\VPKGYSFYLAHEF\LDV
						LPVHKFQKTPQGWREVFVDIDP
						QVSDKLRFVLAPSATPAEAFIQ
İ	}					HDETRDHVEVCPDAGVIIEELS
						QRIALTGGAALVADYGHDGTK
						TDTFRGFCDHKLHDVLIAPG\T
	:					ADLTADVDFSYLRRMAQGKVA
						SLGPIKQHTFLKNMGIDVR/L*R
						FF*INQMRPSVRQQLLQGYDML
				:		MNPKKMGERFNFFALLPHQRL
						QGGRYQRNARQSKPFASVVAG
						FSELAWQ
22411	52779	A	22539	2	377	
22412	52780	A	22540	2	1427	COSPONITH CHOKINGOWN A
22413	52781	Α	22541	2	568	CSSPSN/TILGVQKLNSQWRLA
						QDFRLINEAVIPLYPVVPNPYTL
						LSQIPEEAESFTVLDLKDAFFCIP
						LHSDSQFLFAFEDPTDHNSQLT
						WMVLPQGFRDSPHLFGQAQAQ
		İ				DLVHFSSPGTLVVQYMDDLLL
				i		ATSSEASCQQATLDLLNFLANQ
						GCKRVGIALGVLTQTHGTTPQP
	1		<u>l</u>	<u> </u>		VAYLSKETDVAAK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22414	52782	A	22542	l	1230	GPRGTGPKGKARSEKGCSLSHG
22414	32762	\ \ \	22342		1250	PQTNKPLVVQKGQKMEQANHP
	-					VGLVISVVYKDILKKIVQRETS
						HPLIHVRYAEAITGRRTAPEDK
						GSLGRDMLAKAGAIIYMNMGN
						KLPIWCHLLEEGIYLEVWALEG
						QFGRAKNACPVQIRLKDPTTFP
						YQRQYPLRPEAHKGLQDIVKH
						VKAQGLVKKCSSPCNTPILGVQ
						KPNGQWSLVQDLRLISEAVIPL
		ŀ				YPVVPNPYTLLSQIPEEAEWFP
:						VLDLKDAFFCIPLHYDSHDSQF
		l				LFAFEDPTDHTSQLIWTVLPQG
			}			FRDSPHLFGQALAQDLGHFSSP
						GTLVLQYVDDLLLATSSEASCQ
						QATLDLLNFLANQGYKASRSK
						AQLCLQQVKYLGLILARGTRTL
1		1	ļ			GKERIQPILAYPHPKTLKQLWG
						FLQIT/GFCQLWIPR*SKI
22415	52783	A	22543	1	211	i sąm or ogswark ora
22416	52784	A	22544	1	472	
22417	52785	A	22545	1	393	
22418	52786	A	22546	59	411	SWPSDKQTLVVQRGQKMEQA
						NHPDPTDHMSQLMWT/VLPQG
						FRDSPHLFGQALAQDLGHFSSP
						GTLVLQYVDDLLLATSSEASCQ
						QATLALLNFLANQGYK/LSRSK
						AQLCLQQVKYLCL
22419	52787	Α	22547	43	236	TSSLLRKEDSAPSQGKSVVFTL
		1				TSQG*VPLWRTLQLQGPFFTPSS
						RK*LELSSANSQQHLGCPV
22420	52788	Α	22548	1	1446	
22421	52789	Α	22549	1	1521	
22422	52790	Α	22550	2	689	
22423	52791	Α	22551	1	64	
22424	52792	Α	22552	636	1586	SNDRTEDDCGKHPFMSSPP\TEP
						WVCLIIEGQEIDFLLDTGTTFSV
						LIPCLGRLSSRSVTIQGILGQPVT
						RYFSHLLSCNWETLLFSHAFLV
		l				MPESPTPLLGRDILAKAGAIISM
						KTGNKLPICCPLLEGINPEVWA
						LEGQFGRAKNAHPLQIRLKDPI
						SFPYQRQYPLRPEAHKGLQDIV
						KHLKAQDSVRKCSSPCNTPILG
						VQKLNSQWRLVQDLRLINEAVI
						PLYPVVRNPYTLLSQVPEEAEW
						FTVLDLKDAFFCIPLHSDSQFLF
						AFEDPTDHTSQLMWMVLPQRF
		<u> </u>				RDSPHLFGQAQAQDLGHFSSPG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22425	52793	Α	22553	292	1493	SNDRTEDDCGKHPFMSSPP\TEP
		l				WVCLIIEGQEIDFLLDTGTTFSV
						LIPCLGRLSSRSVTIQGILGQPVT
1						RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
		İ				KTGNKLPICCPLLEGINPEVWA
		l				LEGQFGRAKNAHPLQIRLKDPI
						SFPYQRQYPLRPEAHKGLQDIV
						KHLKAQDSVRKCSSPCNTPILG
		ŀ				VQKLNSQWRLVQDLRLINEAVI
						PLYPVVRNPYTLLSQVPEEAEW
						FTVLDLKDAFFCIPLHSDSQFLF
						AFEDPTDHTSQLMWMVLPQRF
		l				RDSPHLFGQAQAQDLGHFSSPG
		1				TLVLQYSEIAKTLYTLIKEMER
						ANTHLVEWEPEAETAFETLKQ
		l				ALVQAPALSLPTGQNFALYVIE
						RAGIALGVLTQTHRTTPQPVAY
22426	52794	Α	22554	2	307	
22427	52795	Α	22555	3	400	
22428	52796	Α	22556	2	78	
22429	52797	A	22557	395	1336	SNNRTEGASSCH\PLTEPQVCLT
						IEGQEIDFLLDTGTAFSVLISCPA
						WGTGQDSVSKKNEKKRKEKA
						KGNHQANAEAKIAARWNPSLE
						IPTEGPLVWNNPLQEIKPQYSLT
						ETEWGLSQGHSFLPLGWLMTE
						EVKVLIPEASQWKILKTLHQTF
				Ī	:	QMGIENTQQMAKSLFTGPNLL
						WTIRHVVKACEVCQRNNPLVH
						HKAPLGEQRIGHYPGEDWQLD
						FTHMPKSKGFQYLLVCVDTFT
						NWIEAFPCKIEKAQEVNKVLIH
		1				ETIPRFGLPQSLQSDNDPAFKA
					;	MITQGIFQGAKDTISPSLCLEATI
						LREGREGN
22430	52798	Α	22558	244	638	GPLNPTFLGPPCGLGGKLVFLL
						LRQ*VQLFRSAGSRDRCGFLDR
						ERNKTNQNCGKQAKKSPKDHK
		l				NPRAVGYVPFNL*QEGNLAQP
						GYMSPSPSLI*SRSRQTWGSFQ
						MILIGTKRNRPKRKSEIRERLQP
22431	52799	Α	22559	210	600	RGKRTWRERRNGWSRGFCPLP
		İ				LTGSWLISRESPRLACLGEGLG
						VPGM*DLPHAWPTTLMCPELV
						GSWSH*LQE*SHGPSQ*VLQLL
						RWCVWSLFLLMFGCVWSFFLL
						VSSWSRWLRSEATDLCGECYSS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22432	52800	Α	22560	2746	5211	ETSCPLVRVHYAEAITGRCTAP
						EDEGSLGQKPPARRSNNRTEGA
						WGKRQLMSSP/STEPRVCLTIEG
		ł				QEIDFLLDTGAAFSVLISCPGRL
						SSTSVTIQGILGQPVTRYFSHLL
						SCNWETLLFSHAFLLMPERPTG
						LLGRDILAKAGAIIYMNMGNKL
						PICCP\LLVEGINPEVWALEGQF
						GRAKNARPLQIRLKDPTTFPYQ
						RQYPLRPEAHKGLQDIVKHVK
						AQGLVKKCSSPCNTPILGVQKP
						NGQWSLVQDLRLISEAVIPLYP
						VVPNPYTLLSQIPEEAEWSWAL
						TFKDAFFCIPLHSDSQFLFAFED
						PTDHTSQLTGTVLPQGFRDSPH
						LFGQALAQDLGHFSNPGTLVLQ
						YVDDLLLATSSEASCEQATPDL
						LNFLTNQGYKVSRLKAQLCLQ
						EVKYLGLILARGTRALSKE*IQP
						ILAYPHPKTLKQLRGFLEITGFC
						*LWIPGYSKIARPLYTLIKETQR
						ANTHLVEWEPEAETTFKTLKQ
	}					ALVQAPALSLPTGQNFSLYVRE
						RARIALGVLTQTHGTTPQPVAY
						LSKEIDVVAKGWPHCLRVVAA
						VAVLASEAIKIIQGKDLTV*TTY
						DVNGILGAKGSLWLSDNCLLR
						YQALLLEGPVLQICTCMALNPA
	1	1				TFLPEDGEPIKHDCQQIIVQTYA
						ARDDLLEVPLTNPDLNLYTDGS
						SFVENGI/RKVSDVTILESKPLPP
						GTSAQLAELVALTWALELGKG
22433	52801	Α	22561	903	4440	DHPGHIHSSPCHERPNNVSTLPF
		-				LPGEQHPILLPPRNCPGDKILEE
						NFRYNNYKRTMMSFKERLENT
						VERCAHINGNRPRQSRGFGELL
						STAKQDLVLEEQSPSSSNSLENS
						LVKDYIHYNGDFNAKSVNGCV
						PSPSDAKSISSEDDLRNPDSPSS
	1					NELIHYRPRTFNVGDLVWGQIK
		ļ				GLTSWPGKLVREDDVHNSCQQ
						SPEEGKVEPEKLKTLTEGLEAY
						SRVRKRNRNGIKVPPRSPRSTIG
						SPRPSMPSSPS
22434	52802	A	22562	1	501	0.11.00.0
44734	22002	/ L	22302		301	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22435	52803	Α	22563	1688	2038	SATHGLSDSCYWLVVSRAIGRN
						NLSPASPASIIHALVFLAEPQLF
						QPRILGLIDEESGHEGGRPAAKT
						RKLSGPWSRPAAFMSTLLINQP
						QYAWLKELGLREENEGVYNGS
						WGGRGEVITTYCPANNEPIARV
						RQA\SVADYEETVKKAREAWKI
		-				WADIP\APKPKEKLVRQIG\DAL
						REEDSKLLRKPWCPLEDWGKIL
					1	LEGVGEVQEYVDICDYAVGLS
		Ì				RMIGGPILPSERSGHALIEQWNP
						VGLVGIITAFNFPVAVYGWNN
		1				AIAMICGNVCLWKGAPTTSLIS
						VAVTKIIAKVLE\DNK\LPGAICS
						LTCGGADIGTAMAKDERVNLL
						SFTGSTQVGKQVGLMVQERFG\
						RHYQNWRSLLELGGNNAIIAFE
					-	DADLSLVVPSALFACCGEQLGQ
						RCTT/SRGRLFYYMESIHDEVFK
		1			}	QTLKRAYGTRSRVG\NPW\DPN
					}	VLYGAISHQGRQLSMFLGAVEE
						AKKERLAQWVYGGKVMDRPG
:						N\YVEPTNVTS\LDHDASIAHTE
						TFAP\ILYVFK\FKNEEEVFAWE*
į						WK*NRGLSS*HSLPKDLGAESF
	1					RW\LGP\KGTQTCG\IVKCQPFPT
						KWGLRLGGA\FGGEKAHLVVA
						RESGQ*C/WGKQYMRKVYLVL
		l				FNYS\KDLPSGPQG\IKFQ
22436	52804	В	22564	1	1434	
22437	52805	В	22565	1	1185	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22438	52806	A	22566	11	2450	GQACHASSSPLKGSRGSPPNFEI
		1				LYGPIFEDSLAPD*PKVIAGQ\IT
		1				LRDAVTGPISSPNEAGKIYQLKP
		1				NPAVLICRVRGLHLPEKHVTW
						RGEAIPGSLFDFALYFFHNYQA
						LLAKGSGPYFYLPKTQSWQEA
		ł				AWWSEVFSYAEDRFNLPRGTIK
		Ì				ATLLIETLPAVFQMDEILHALR
						DHIVGLNCGRWDYIFSYIKTLK
						NYPDRVLPDRQAVTMDKPFLN
		į				AYSRLLIKTCHKRGAFAMGGM
						AAFIPSKDEEHNNQVLNKVKA
	1					DKSLEANNGHDGTWIAHPGLA
						DTAMAVFNDILGSRKNQLEVM
						REQHAPITADQLLAPCDGERTE
	İ					RKFNALMESEKGQSQNLWKFA
ŀ						VYSGLRHGELAALAWEDVDLE
						KGIVNVRRNLTILDMFGPPKTN
						AGIRTVTLLQPALEALKEQYKL
						TGHHRKSEITFYHREYGRTEKQ
İ						KLHFVFMPRVCNGKQKPYYSV
						-
						SSLGARWNAAVKRAGIRRRNP
						YHTRHTFACWLLTAGANPAFIA
						SQMGHETAQMVYEIYGMWID
						DMSDEQIAMLNARVMVYDVA
						SGNALFISELGPLPENVTWLSPE
ļ					:	GEFQKWNGTAWVKDTEAEKL
						FRIREAEETKNNLMQVASEHIA
i i						PLQDAADLEIATEEEISLLEAW
						KKYRSIPLCSKVVTMMFDDRP
						GCPVGFPMWGRGGFDRMPPG
22.422	50005	4.	22565	·	0.41	WGGRPMPPSRRDMSPHRGPPPP
22439	52807	Α	22567	1	941	MIPNVQGPMRGLEGPNYIFIKIS
						KNRRYTMAVTSVGLASSLCRV
						VRPCRAPAGVVSRGSPDDEWL
						FCPQARIQQQQDID/NGTLPDFIS
						ETASIRDA/DWKIRGIPADLEDR
						RV/EITGPVERKMVINALNAI\F
						MADFEDSLAPDW/NKVIDGQIN
						LRDAVNG/TISYTNEAGKIYQL
		1				KPN/PAVLICRVRGLHLPEKH/V
		1				TWRGEAIPGSLFDLA/LYFFHNY
						QALLAKGSGP/YFYLPKTQSWQ
						EAAWWS/EVFSYAEDRFNLPRG
						T/IKATLLD*TLAGGFQMVDPSA
						LGG*TGPWDHLRYIKVENIPKV
						GEEGDAETTGGTAV
22440	52808	A	22568	2	393	
22441	52809	Α	22569	1066	1824	
22442	52810	A	22570	2	165	
22443	52811	Α	22571	3	902	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22444	52812	Α	22572	1	2034	
22445	52813	A	22573	176	873	RLFGEMSKLN/HNLSEVMSKLE KQHSSKVFVVKRLSSSSRRSLVI SPPVRTSTVSSPLTSPTSPSTLSS KSKSESVSATEDLAPDAAQGED NSEIKELLEDEEIGKEGSEASSS EEEEPLLTCNGPSQAQPSPIIESA RSQEEVPPSSPALSPGGALSPSG HPLSSAAEVVLRTRTAVPQAKG NNVVITSYMTNRGFFEDKKATF APSFLMNIKGNKTSVVKNSILE QGQLTVN
22446	52814	A	22574	1	1890	QGQLIVN
22447	52815	A	22575	2	1279	LSVFPRVRNFTTQAECH/KLFK VLKNFLIAGGGVHESSKRAS*T LQFIYY\SEWYGHEELKAIVWN NDLLWEDYEEKLADQAVRTM EIYVAQFSEIKERIAKRGRKLVD YDSARHHLEAVQNATKKDEAK TAKAEEEFNKAQTVFEDLNQEL LEELPILYNSRIGCYVTIFQNISN LRDVFYREMSKLNHNLYEVMS KLEKQHSNKVFVVKGLSSPSTL SLKSESESVSATEDLAPDAAQG EDNSEIKELLEEEEIEKEGSEASS SEEDEPLPACNGPAQAQPSPTT ERAKSQEEVLPSSTTPSPGGALS PSGQPSSSATEVVLRTRTASEGS EQPKKRASIQRTSAPPKPPEKPV RTPEAKENENIHNQNPEELCTSP TLMTSQVASEPGEAKKMEDKE KDNKLISANSSEKYIFQRPSVLI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22448	52816	Α	22576	1	1791	MPGARTSSSGASENHRARGQG
		ŀ				GGPQGVGRMAEGKAGGAAGL
						FAKQVQKKFSRAQEKAL*KLG
]	KAVETKDERFEQSASNFYQQQ
						AEGHKLYKDLKNFLSAVKVMH
						ESSKRVSETLQEIYSSEWDGHE
		Ì				ELKAIVWNNDLLWEDYEEKL
		ŀ				ADQAVRTMEIYVAQFSEIKE\RI
}						AKRGRKLVDYDSARHHLEAVQ
		1		ļ		NAKKKDEAKTAKAEEEFNKAQ
					İ	TVFEDLNQELLEELPILYNSRIG
						CYVTIFQNISNLRDVFYREMSK
						LNHNLYEVMSKLEKQHSNKVF
						VVKGLSSSSRRSLVISPPVRTAT
						VSSPLTSPTSPSTLSLKSESESVS
		ŀ		:		ATEDLAPDAAQGEDNSEIKELL
						EEEEIEKEGSEASSSEEDEPLPA
		}				CNGPAQAQPSPTTERAK\SQEE
		1				VLPSSTTPSPGGALSPSGQLSSS
		1				ATEVVLRTRTASEGSE\RPKKT
	}	1				ASIQRTSAPPRRPPPPRATASPRP
		1				CSGNIPSSPTASGGGSA\TSPRAS
		1				LGTGTASPRTSLEI*PNPE\PPEK
						PVRTPEAKENENIHNQNPEELC
		1				TSPTLMTSQVASEPGEAKKME
						DKEKDNKLISADSSEGQDQLQV
			<u> </u>			SMVPENNNLTAPEPQEEVSTSE
22449	52817	Α	22577	2216	2568	RPSLGLGM/C/TIWQ/TSLA/GKH
						RRTIETHTDDYRISLEQGTPGLE
						PPLNALTAISKKIGCNGFFPFQIR
						PGKNETDGRMFSPAIGIVEDPV
		1				TGNANGPMGAWLVHHNVLPH
						DGNVLRVKGH
22450	52818	A	22578	108	532	WGSKGNLLYSHQTHSVSHQFS
						LVACCLELEYSPFSFPYACLLSS
						LMRLEKHQELNLWPAHQRSMR
						HPQDTLVVFS/HHGRVLL*DKT
						SRGTIRQQHSRFTKICCSAATAA
						VTQANRVWSGPLANSNRPAAE
						GPVLAAFMGWL
22451	52819	Α	22579	1	1125	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	10000	 		<u> </u>	loos.	
22452	52820	Α	22580	I	985	MCARGSLWGCYSGLERKCEAA
						LPDRVVSSKSACGAVSQDLLQ
						AQGRRAGPPSWLYADPVHTGT
		1				IECGGHWCLNAALLDFQSPQRL
	[1				LTYLHCGLQRPMESDPPELAPM VHHQEGIYLNIRKAMYDKPTA
		İ				NKLNWDKSNVSSLITGTRQEET
						TANGTTFATIHLQFFIQIPAVPR
				•		KTLLNRADYYFPEHNPSLTLVI
:		1				NRRITKFSEPNWEAPPSRGRLTP
;						HTAGYSSETKIPEERSDSSIRGS
		1				RKSAVLQPLLLDSQANRVWSG
		1				PLANSNRPAAEGPVC*LWWVT
		1				LHGPLKATMQVCQQPLW
22453	52821	A	22581	294	934	YPPSPAELTVCRLRQLQLRDVL
						YCIAYPSGPGVKPFRGNSA/G/V
						VFPADNLSEAQMQLI/ARELGH
		1				SETAFLLHSD/DSDVRI/RFTPTV
		1				EVPI/CGHATVAAHYVRAKVL/
		Ì				GLGNCTIWQTSLAGKHR/VTIE
		Ì				KHNDDYRISLNALTAIS/KKIGC
						NGFFPFQIRPG/KNETDGRMFSP
						AIGIVE/DPVTGNANGPMGAWL
		<u> </u>				VH/HNVLPHDGNVLRVRQ*GPR
22454	52822	В	22582	1163	5585	A G A A A A A COLUMN DAY
22455	52823	Α	22583	2	429	AACAAAMSLVIPEKFQHILRVL
		1				NTNIDGRRKIAFAITASSSSSSSS SSSSSSSSSSSSSSSSSSSSSSSSSSS
		1	1			SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
						VGRRYAHVVLRKADIDLTKRA
						GELTEDEVERVITIMQNPRQYKI
						PDWFL
22456	52824	A	22584	84	424	GVHAASDKPCWITTSQTPDVQ
					}	HRW/WLLISSML*TAPNRRSDK
		1				AFTPHPTNHAGSQLRKHPTCSI
		ł				DGVIDIFHAVNTFVQHHMEETL
						KAFHVTGHHILIARHGLGVGKE
					1	NTEHAANVVNDQRNSGFFCRF
		ł				QQTFCQTCCQLIQRFVNARLRD
						FISC
22457	52825	Α	22585	88	381	IQVLANGLDNKLREDLERLKKI
						RAHRGLRHFWG*VGGLISLPTS
						TQHSSYSLFFFPNLLFLLCMTCD
						SSLFTCSLRVRGQHTKTTGRRG
20172	150625	<u> </u>	22525	255	(00	RTVGVSKKK
22458	52826	Α	22586	355	699	FIDTTEAGSHFLYALVRFAATIT
						TITDDGARRLRTLFQRQQADLV
						SVSKSGFFTADCTHTNALIDVV
			[RTIFNNAVF*NPGFVIARLKIEIA
			}			IIKATFGQLTEDREQVLMFQAV RR
<u> </u>	<u> </u>		<u> </u>		<u></u>	IUN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22459	52827	Α	22587	1389	1532	SVLPGRPDARQYRQTRQSW*R
						WSLRLRGSYPVAGRSLSRRHRQ
		ŀ		ĺ		TERF
22460	52828	Α	22588	1	427	LVHQQWIQTKKKSLIYPFCHSP
						SQRVVEVAALDQWLAMLYPR
						VEHQAVLAVVVVVVGPLVVRP
						YHTRRDAVRLLRQPAVFIVAV
		•				DIVRILRDAVIRPHCECTVRV\R
						SGSGSRLPAGSVAYGMRMSRS
						GARVRTLSILPVAL
22461	52829	Α	22589	1	915	
22462	52830	Α	22590	1166	1321	
22463	52831	Α	22591	813	977	
22464	52832	Α	22592	3485	4237	
22465	52833	Α	22593	1	499	ARAAACAAAMSLVIPEKFQHIL
					1	R\VLNTNIDGRRKIAFAITAIKG
						VGRRYAHVVLRKADIDLTKRA
						GELTEDEVERVITIMQNPT\QYK
						IPDWFLNRQKDVKDGK\FQPGS
						*ANGLSTTKL\RENLKPLKEDSG
						PIKGLRQLP/WGLRVRGQHTKT
22466	52834	A	22594	1	1197	TGRRGRTVGVSKKK
22467	52835	A	22595	1	1905	
22468	52836	A	22596	37	514	WTWWLRCYENRHRHCHLLQQ
22400	152050	``	22330	[]		LPTSPHQHAVATPTTVAVPTAP
						AATPPVVVAVVVAPVVVVVP
						VVPVVASVPFVEPGSDVIVCAE
						MDEQWGYVGAKSRQRWLFYA
						YDSLRKTVVAHVFGERTFDVVI
						WMTDGWP\LYESRPAFTGRDC
		1	ł			VDDAFFAVAGV
22469	52837	A	22597	1	471	
22470	52838	Α	22598	2	323	SRIQPGSDVIVCAEMDEQWGY
						VGAKSRQRWLFYAYDSLRKTV
			l			VAHV/FGERTIRTLTRRLFRCLR
						SSHDLVSSPVSHRRRRAGSRQR
						HRRARGGENVEYQWGALRHG
22471	52839	Α	22599	2033	2310	ARYVGNAGCTGGALFGRSQTI
						REAVG/ERRLAESRDVAALDTP
						CDPEIKMAEWMQTLKETGSTS
						GHIVTRRISVRISHVTPGPASQD
						GPDVQQR
22472	52840	В	22600	1	4891	
22473	52841	В	22601	7	1233	
22474	52842	A	22602	193	492	
22475	52843	Α	22603	1	486	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22476	52844	Α	22604	62	443	RSDRLLIHGFSFPAMGFRLRKT VVAHVFGERTMATLGRLMSLL SPFDVVIWMTDGWPLYESRLK GKLHVISKRY\TQRIERHNLNLR QHLARLGRKSLSFSKSGTACCG HCLTRDIPMLHHQGNQLF
22477	52845	Α	22605	971	1444	SMGSGDMRSSPSRAAALRQLPR PFRQLAVRMEMISRFRSAKEQT KSLRKWRKAKSIF*SVAQTGFN VLVCTTIIETGIDSRQPTLLSLNA RITSVCAAAPVTRSRRTFASSGY AWLLTPHPKAMTTDAKNVLKQ LPRWKISGQVRAGTARSGDSRR W
22478	52846	Α	22606	1	1047	
22479	52847	Α	22607	318	598	LTGFRRLCDLRFGGGAVRSVLA RSREFILQH*FRRIVTGSMGRSL VPSAPLR/PSIVLVLFRKAW*LV KTQRKMHVVSIVQKKASCWYP LLGCL
22480	52848	Α	22608	1	2652	
22481	52849	Α	22609	1	852	
22482	52850	Α	22610	1	807	
22483	52851	Α	22611	1	1740	
22484	52852	A	22612	1	3144	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22.12.5	Lines		00610	1.72	11000	DVOVGCODI CAL DCA DI JEGDOV
22485	52853	Α	22613	173	1983	PVQNGSSPLGALPGAPLIFGPSV
		ļ				PAISRPGSCSPEWPTLDTSGYSA
		ĺ				PR/WLFTAYAGSEAVFAHVFGD
	!				j	ALGDAGLLMSLLSPFDVVIWM
						TDGWPLYESRLKGKLHVISKRY
						TQRIERHNLNLRQHLARLGRKS
						LSFSKSVEQHDKLESLPGQIDKS
						LDEASLSLRAGSLRTITHILLPLL
						RPAILSALIYSFVRAITTVSAIVF
						LVTPDTRVATAYILNRVEDGEY
						GVAIAYGSILIVVMLAIIFIFDWL
						IGESLLSRFEKTVDDHSQSRLPA
						EGSILSQDISGVKLFSLLTPARG
						YVPIRAPCPFLCASYRFFCVARQ
						QGLDAALIDAPLAVDALTPLSE
1						LLSHVGQAPCSVPVVDEDQQY
						VGIISKGMLLRALDREGGVRVP
			İ			VDYILNGFHPLLLGMPAPLAIIV
						FALIACRFPGSNGRIRRSRRIRH
						LHKMPDATLSASYQAYVRHQT
						NVTALVLRSSHRFRKDRHCWG
	ı					FVFLQQPILRATGQFEIQTFRYL
						SVFSVQRHKTNGKTALMSGNV
						ADKAVNRATGPTACVWLFSLR
						NSSMVTTRFCGLCEWADQQGA
						RPARISENGKRGYSYLYDVHM
		1				GAQSGETRGRQGEYASKPPSRE
22486	52854	В	22614	1	2149	
22487	52855	A	22615	1	1815	
22488	52856	Α	22616	2014	3411	
22489	52857	Α	22617	1	922	MVICQPINGGWQWPPSTGTDP
		1				APYFRIFNPTTQGEKFDHEGEFI
						RQWLPELRDVPGKVVHEPWK
						WAQKAAAPPQTRLHSTPTTIPH
					·	RVPGHTPNLKPIWEVDLPKFSH
						KATSHATDKPSDTTQNDQIVIA
						YMEPPEIVHETTTSSTHPSLYIIL
						KRQTRELRPQSVTSRIQPGSDVI
						VCAEMDEQWGYVGAKSRQRW
						LFYAYDSLRKTVVAHVFGERT
						MATLGRLMSLLSPFDVVIWMT
						DGWPLYESRLKGKLHVISKRYT
		1				QRIERHNLNLRQHLARLGRKSL
		i				SFSKSVELHDKVIGH\YLNIKHY
L	<u> </u>	L	L			OF SNOVEL FIDE VIOLATENIKHY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ł	09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22490	52858	Α	22618	1458	2185	YGHEWRWMPGNRPH\YGRWP
					,	QHDFPPFKKLRPQSVTSRIQPGS
		ŀ				DVIVCAEMDEQWGYVGAKSR
						QHWLFYAYDRLRKTVVAHVF\
						GERTMATLGRLMSLLSPFDVVI
						WMTDGWPLYESRLKGKLHVIS
						KRYTQRIERHNLNLRQHLARLG
						RKSLSFSKSVEKITSGPTTNATP
						GTQKTATDKRGGKKLPIQIPIGP
						ISRIASGAAINTISIGLRKFLVTA
						GVMLSTQRSIYDRHQVITSAGIT
22491	52859	Α	22619	493	1518	
22492	52860	Α	22620	963	1597	
22493	52861	Α	22621	1	3052	MATDSVGLNCTPAPHARDVLL
						WRPSSLPVSTSCVFPLGRIGDVS
		l		:		DESDDKTEAPTPHRLEKAREEG
						QIPRSRELTSLLILLVGVSVIWF
						GGVSLARRLSGMLSAGLHFDH
						SIINDPNLILGQIILLIREAMLAL
						LPLISGVVLVALISPVMLGGLVF
						SGKSLQPKFSKLNPLPGIKRMFS
						AQTGAELLKAILKTILVGSVTG
			•			FFLWHHWPQMMRLMAESPITA
						MGNAMDLVGLCALLVVLGVIP
						MVGFDVFFQIF
22494	52862	Α	22622] 1	1233	MANDLPHSNSGDGQKPSSPDLF
		Ì				TAPLLADPTVGFLPNDAEELFIF
						LTEITEITIPCRVTDPQLVVTLHE
		ŀ				KKGDVALPVPYDHQRGFSGIFE
		ļ				DRSYICKTTIGDREVDSDAYYV
						YRLQAVPPVQLLTGWCVTAKA
						PPDISAISALTAVKHGNCSSLTP
ŀ		ŀ				LLNPPGSDVIVCAEMDEQWGY
•						VGAKSRQRWLFYAY\DSLRAR
1						RVVAHVFG*RTMATLGRLMSL
		ľ				LSPFDVVIWMTDGWPLYESRL
						KGKLHVISKRYTQRI\ERHNLNL
						RQHLARLGRKSLSFSKSVELHD
						KPSPQHREGVKDVMRFNEPMY
						PLESRPRLYVCHLRNGAQASGR
						RCQLPMGCACHPVSQRNDRPN
						RRPERHATRKTTTRQRQQNSRQ
						HKQTQKKAKPNKRRATERGKG
	<u> </u>	<u> </u>	<u> </u>	l	<u> </u>	PQQKAERPKRHTHCFLKLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22495	52863	I A	22623	3	1386	HEASSPLAMASLTVKAYLLGK
22493	32803	A	22023]3	1300	EDAAREIRRFSFCCSPEPEAEAE
						1
				•		AAAGPGPCERLLSRVAALFPAL
·						RPGGFQAHYRDEDGDLVAFSS
·						DEELTMAMSYVKDDIFRI*IKE
	1				•	KKECRRDHRPP\CAQEAPRNMV
						HPNVICDGCNGPVVGTRYKCS
						VCPYY\DLCSVCEGKGLHRGHT
				•		KLAFP\SPFGHL\SEGF\SHSRWL\
						RKV\KHGHF\GWPGW\EMGPPG\
, ·						NWEPTFLPRAG\EARPGP\TAES
						ASGPSEDPSVNFLK\NVGESVA\
						AAFSPL\NFEVDIDVEHGGKRSR
						LTPVSPESSSTEEKS\SSQPSSCC
					•	SDPSKPGGNVEGATQSLAEQM
			·			RKIALESEGRPEEQMESDNCSG
	1.	ľ				GDDDWTHLSSKEVDPSTGELQ
						SLQMPESEGPSSLDPSQGGNPQ
						GWKEAALYPHLP\PEVEPAA*F
						ESLSQMVVPWGFS*LKGGWVH
•						QGSCRPRNYDI\GAAL\DTIPVIQ
22496	52864	Α	22624	1032	1127	FYAKAEAGDRTVA*PHKPKMF
						LSIHHRVAKI
22497	52865	A	22625	121	561	VFPGPLGSLVDICGPSRVPFPWV
		_				VPAVLPVQFVQRLPRFFRCPPV
						LVTSVIRSISTLIQALGSSPFARR
	1 .]			YWGNLG*FLFLGVLRCFSSTGS
	Ì		ļ			PH*PMNSVNDSVSKHT/EVTPF
						GN\PGNNGSYHLTDAYRRLARP
1	1					SSPLT\PGHPPLYG
22498	52866	A	22626	243	454	ATAGCPSCSATEGVVRNGKST
22498	32000	^	22020	243	434	I
						AGHQRYLCSPCRKTWQLQFTY
		ĺ				TASQP/GARIMGVGLNTVLRHL
22.425	500.5	ļ.	22/25			KNSGRSR
22499	52867	A	22627	3	640	
22500	52868	A_	22628	236	410	
22501	52869	Α	22629	116	359	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
l	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22502	52870	A	22630	276	1905	LFLYSSHGQYGCPSCSATDGVV
						RNGKSTAGHQRYLCSHCRKTW
						QLQFTYTASQPAGQ*RTAIATVI
						CCNNCQRRHTNSRCHTDNCRR
						ADSASSHATSGSSSRGCAGSGR
						CGASSTRRCLGPVCCFRGSKRS
						WRRQHEGHKDCRKLVFHNFSFI
						HRMDNTAVPPVQLLTGWCVTA
						KAPPDISAISALTAVKHGNCSSL
						TPLLNPPGSDVIVCAEMDEQW
						GYVGAKSRQRWLFYAYDSLRK
						TVVAHVFGERTMATLGRLMSL
1						LSPFDVVIWMTDGWPLYESRL
1						KGKLHVISKRYTQRIERHNLNL
						RQHLARLGRKSLSFSKSVELHD
						KQQGLDAALIDAPLAVDAQTP
						LSELLSHVGQAPCAVPVVDED
<u> </u>						QQYVGIISKGMLLRALDREGGV
				·		RVPVIYPKRFRTTDVCMPLSLG
						TCIREIEAEFPDPIFRRSVVYICE
						HNTNGAMGIIVNKPLENLKIEGI
		1				LEKLKITPEPRDESIRLDKPVML
						GGPLAEDRGFILHTPASNFASSI
						RISDNTVMTTSRDVLETLGTDK
						QPSDVLVALGYASWEKGQLEQ
						EILDNAWLTAPADLNILFKTPIA
						DRWREAAKLIGVDILTMPAQD
						GTPDWNIIERLLKEWQPDEIIVG
						LPLNMDGTEQPLTARARKFAN
						RIHGRFGVEVKLHDERLSTVEA
1						RSGLFEQGGYRALNKGKVDSA
						SAVIILESYFEQGY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22503	52871	Α	22631	399	2337	PEVQGLQQAGAAARRPGVGLA
22303	32071	` `	22031		2337	APRVVCEGRLRGSGLQSRLCPG
						ASV*ATDGVVRNGKSTAGHQR
						YLCSHCRKTWQLQFTYTASQP
		1				AGQ*REKEEKEEVEEEEEEKK
						KRREGEKEEKEKEKKKK
						FKKKKKKKKKKKKKKKKK
						KKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKKK
						WRNEARTTGACLPSPTVSAAAP
İ						PIVSTTVTAFAAATLAKGDLRS
						RASSKQVPLLVVLVWVWPHPG
						WFVKAGSVALGSSPGCVLELL
						CKLLTGWCVTAKAPPDISAISA
						LTAVKHGNCSSLTPLLNPPGSD
						VIVCAEMDEQWGYVGAKSRQ
						RWLFYAYDSLRKTVVAHVFGE
						RTMATLGRLMSLLSPFDVVIW
ĺ						MTDGWPLYESRLKGKLHTQER
						KARSKQQKDADTAPPIQVGTQ
						TPQTVHAARDTRQASPRASKIS
						QHRQADGDKRQTRHAEAQAT
		l				RTHTGRRKTIKETATQQAYADE
ł						QTRRTARTHRRTARHTAAHTH
		İ				RTRVAARQTTEKDNEQSDTQPP
	ľ					TQKTPSATATAHTPRRTETTTNI
						VQARHTSARERNHTRQHQATR
						RETTINAHTNTPHYHAEHTQSR
						SQQQTTSTERPTTDSSTTINNNN
		1				HTPRTVRDERRMTQHKHPSRA
						STTTNRTTTTGAYQHTQRDTRD
		l				TRRDTHEDTTTQQPHTTTPTAN
		<u>L</u>				EAAYTAHTTKTDDHRHTADDT
22504	52872	A	22632	103	313	ATAGCPSCSATEGVVRNGKST
			}			AGHQRYLCSPCRKTWQLQFTY
		1				TASQP/GARIMGVGLNTVLRHL
			L			KNSGRSR
22505	52873	A	22633	620	832	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	deterion, \=possible nucleotide insertion)
				Sequence		
22506	52874	Α	22634	453	1047	AADDRGHLRPTDGLLRNGKST
						AGHQRYLCSHCRKTWQLQFTY
						TASQPAGQ*RVTGTSERREQIIQ
						RLRQQGSVQVNDLSALYGVST
ļ						VTIRNDLAFLEKQGIAVRAYGG
						ALICDSTTPSVEPSVEDKSALNT
						AMKRSVAKAAVELIQPGHRVIL
						DSGTTTFEIARLMRKHTDVIAM
						TNGMNVANALLEAEGVELLMT
						AGICALLTGCCVTAKAPPDISAI
						SALTAVKHGNCSSLTPLLNPPG
						SDVPVCAEMDEQWGYVGAKS
						RQRWLFYAYDSLRKTVVAHVF
						GERTMATLGRLMSLLSPFDVVI
ŀ						WMTDGWPLYESRLKGKLHVIS
					:	KRYTQRIERHNLNLRQHLARLG
						RKSLSFSKSVELHDKVIGHYLN
						GDFLQSTLREAPLIPHTREHP
22507	52875	Α	22635	1089	1686	TSPGYWRANSTDPDGRYCRTG
						TSGGVCHALSSLRCPSCSATDG
1					1	VVRNGKSTAGHQRYLCSHCRK
						TWQLQFTYTASQPAGQ*RFLSK
						QEKTLLINQLASEEQQKVTSRL
						SDALRNGRVWQLAIIYLTIQVA
						VYGLIFFLPTQVAALLGTKVGF
						TASVVTAIPWVAALFGTWLIPR
						YSDKTGERRNVAALTLLAAGIG
						IGLSGLLSPVMAIVALCVAAIGF
				1		IAVQPVFWTMPTQLLSGTALAA
						GIGFVNLFGAVGGFIAPILRVKA
						ETLFASDAAGLLTLAAVAVIDR
						RGTSLNRLRNTMIALLSNPFKE
						RLRKGEVQIGLWLSSTTAYMA
						EIAATSGYDWLLIDGEHAPNTI
						QDLYHQLQAVAPYASQPVIRPV
						EGSKPLIKQVLDIGAQTLLIPMV
						DTAEQARQVVSATRYPPYAVH
i						PVQLLTGWCVTAKAPPDISAIS
	:					ALTAVKHGNCSSLTPLLNPPGS
						DVIVCAEMDEQWGYVGAKSR
						QRWLFYAYDSLRKTVVAHVFG
						ERTMATLGRLMSLLSPFDVVIW
						MTDGWPLYESRLKGKLHVISK
						RYTQRIERHNLNLRQHTGTGW
22500	53076	_	22626	01	200	DGSRCRSQKSGGAA
22508	52876	Α	22636	91	200	SGIIAFQYLLMTLKVLPVASVSI
						SCPS\CSATDGVVRNGKSTAGH
						QRYLCSHCRKTWQLQFTYTAS
	<u> </u>	L	<u> </u>	l		QPAGQ*RGGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22509	52877	Α	22637	2649	3270	GAPVASVSISCPSCSATDGVVR
						NGKSTAGHQRYLCSHCRKTWQ
						LQFTYTASQPAGQ*RTIGRAML
						KNKHMTITEFVAQRMRDAQQK
						GEISPDINTAMTSRLLLDLTYGV
						LADIEAEDLARKRRLLRITRDD
						WRYLNRILILSLFRRAGDNCAR
						VSYQTPTREIKKLDGLWAFSLD
ŀ						RENCGIDQRWWESALQESRAIA
						VPGSFNDQFADADIRNYAGNV
						WYQREVFIPKGWAGQRIVLRF
						DAVTHYGKVWVNNQEV
22510	52878	Α	22638	1	2470	MNADTDYSIAEAAFNKGETAM
}						TINGPWAWSNIDTSKVNYGVT
						VLPTFKGQPSKPFVGVLSAGIN
ŀ						AASPNKELAKEFLENYLLTDEG
						LEAVNKDKPLGAVALKSYEEE
						LAKDPRIAATMENAQKGEIMP
1						NIPQMSAFWYAVRTAVINAAS
						GROTVDEALKDAOTRITNCCRI
						RRESLIRPGMRKNPMDVIKKKH
			}			WWQSDALKWSVLGLLGLLVG
						YLVVLMYAQGEYLFAITTLILS
1						SAGLYIFANRKAYAWRYVYPG
						MAGMGLFVLFPLVCTIAIGFNN
						YRRTKQVAGMPAKSSSLRSSQ
						DREMRPGAKAERTRYDSPGSY
						GVAGPGFLGSRGGSPFGGPGKY
						NNTITKYALNIAPSAQNQSPHG
			i			PKLNRRQETQHALDWLDAGYT
						FTYSEDQALTPYFKLAYVYDDS
						NNDNDVNGDSIDNGTEGSAVR
						VGLGTQFSFTKNFSAYTDANYL
						GGGDVDQDWSANDLTGITAKD
						AQMLSVVKPLQEFGKLDKCLS
						RYGTRFEFNNEKQVIFSSDVNN
						EDTFVILEGVISLRREENVLIGIT
						QAPYIMGLADGLMKNDIPYKLI
						SEGNCTGYHLPAKQTITLIEQN
						QLWRDAFYWLAWHNRILELRD
						VQLIGHNSYEQFAHIIFMIAAM
				ŀ		QMGSVWGDDFSSGQHSVARKS
						QLHKPILRVKCHEGFTISANPNS
						SPRYPTHEQAEQPRIVDFSIVRF
22511	52879	Α	22639	1	852	
	1	ш	L	l	<u> </u>	

SEQ ID	1		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		0,7,0,10,21,	sequence	or pepular request	, , , , , , , , , , , , , , , , , , , ,
22512	52880	Α	22640	1	747	MIIIDPRYTDTGAGREDEWIPIR
						PGTDAALVNGLAYVMITENLV
	:					DQAFLDKYCVGYDEKTLPASA
						PKNGHYKAYILATD/GVVRTAK
						STAGHQRYLCSHCRKTWQLQF
						TYTASQPAGQ*RYTDTGAGRE
						DEWIPIRPGTDAALVNGLAYV
						MITENLVDQAFLDKYCVGYDE
						KTLPASAPKNGHYKAYILATDG
						WCVPPKAPPDISAISALTAVKH
						GNCSSLTPLLNPPGSDVIVCAE
						MDEQWGYVGAKSRQRWLFYA
			:	1		YDSLRKTVVAHVFGERTMATL
						GRLMSLLSPFDVVIWMTDGWP
						LYESRLKGKLHVISKRYTQRIE
						RHNLNLRQHLARLGRKSLSFSK
						SVELHDKVIGHYLNIKHYQ
22513	52881	A	22641	1722	2463	NPRLAGSEGSCPSCSATDGVVR
						NGKSTAGHQRYLCSHCRKTWQ
						LQFTYTASQPAGQ*RKLTPGCL
						FVALKGERFDAHDFADQAKAG
		1				GAGALLVSRPLDIDLPQLIVKD
						TRLAFGELAAWVRQQVPARVV
		1				ALTGSSGKTSVKEMTAAILSQC
						GNTLYTAGNLNNDIGVPMTLL
						RLTPEYDYAVIELGANHQGEIA
						WTVTHPPRFVTKQWLCFLPDG
		1				LLLFAEARFVIWQMQPAVMAV
		1				DVAAFAHLFNQQQYQRFTGRP
						PDSSKLPWKITCNGECVTFHW
		l				WLLGVAVRISRLLPVRVMTAL
						EGNYNGTEGISALPFNGIILAHS
		1				NESEWVTFRNNKNNEAFLDRV
		1				YIVKVPYCLRISEEIKIYEKLLN
		1				HIFECWQHFLNNPESLQCYIWA
}		ł				ANADFREARQVIFKTFFTVGVE
		[EELGIRQTRTNDFLVTGDNLLRI
		}				FRFDVGNEDKVRQQFAVVRIH
		1				REVFLVTFHGVNQRFSRHREEF
		l				LFEFCVSVTANLTRCFFGIGFNL
						RLTFFVIGDDFPTLTQNFRILIGV
						IDGEFRLAHKAVAANHAIGLNA
						QNGCRNDFVAHAAGSRALAAD
		1				GRSLEHAPTVETVVAGPLCESG
						DVFTQQEGGNVETRALPEVKA
						AVPPVQLLTGWCVTAKALPDIS
						AISALTAVKHGNCSSLTPLLNPP
						GSDVIVCAEMDEQWGYVGAKS
L		<u> </u>	L			RQRWLFYAYDRLRKTVVAHRL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i .	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>	sequence		
22514	52882	Α	22642	90	1254	KGLCSHSEMVVPSYSATDGVV
						RNGKSTAGHQRYLCSHCRKTW
						QLQFTYTASQPAGQ*RDIVLTK
						LMMVWHSQSGYEKAYAHIQR
						WWYRAILLLTGWCVTAKALPD
						ISAISALTAVKHGNCSSLTPLLN
						PPGSDVIVCAEMDEQWGYVGA
						KSRQRWLFYAYDRLRKTVVAH
						VFGERTMATLGRLMSLLSPFDV
						VIWMTDGWPLYESRLKGKLHV
						ISKRYTQRIERHNLNLSSTSILVL
		Ì				LNGATQFQDLGDLNKELEETY
						TDSKNHISCVGNDFGFNDIFSR
						YVEAVGREGDVLLGISTSGNSA
						NVIKAIAAAREKGMKVITLTGK
						DGGKMAGTADIEIRVPHFGYA
						DRIQEIHIKVIHILIQLIEKEMVN
						FTGLVQRGGGTGPHKDGWGIT
						FYEVAFEDVAVNFTQEEWALL
						DPWQKKLYRDVMLETYRNLAS
						VGDDDNIPSLREQVAHQRYFKT
						WHVEREYFSK
22515	52883	Α	22643	690	1233	RLQSLHYQGRGKPVPMRSYSLS
						FLFTISNSAGVFRCPSCSATDGV
						VRNGKSTAGHQRYLCSHCRKT
ł	ŀ					WQLQFTYTASQPAGQ*RLLGP
						KNRPREQNWLLWCERGVMSRL
						QALYLREQGFNNVKVDNFQTK
						VDSCTRPLAGQQFYPCIPDDAL
	1					VQPLFGGNFVFQRREAGHITNR
						EDIVRHQHGGWRGANSRDDTP
						GTVELFNGFNQQSAIAKTLRAF
						NATRQHDHVIFAISHFDQRRIR
						QQLHAARAGNRQVTIAGDAGS
ŀ				<u> </u>		GDFDTAANQQIDSGDGFSLFTT
						RGEANQCRCAHIHSPFFLPFQT
				1		VRVFSAVPPVQLLTGWCVTAK
						APPDISAISALTAVKHGNCSSLT
						PLLNPPGSDVIVCAEMDEQWG
						YVGAKSRQRWLFYAYDSLRKT
			1			VVAHVFAVKPYWDTGGGFHC
						RWIYMQQEKELLQLSLQQGKY
]			NQKRAAELLGLTYHQFRALLK
	1			· · · · · · · · · · · · · · · · · · ·	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22516	52884	A	22644	1314	1483	RIQRYCGKTGCPSCSATDGVVR NGKSTAGHQRYLCSHCRKTWQ LQFT\YTASQPAGQ*RMFNADG SEVAQCGNGARCFARFVRLKG LTNKRDIRVSTANGRMVLTVT DDDLVRVNMGEPNFEPSAVPF RANKAEKTYIMRAAEQTILCGV VSMGNPHCVIQVDDVDTAAVE TLGPVLESHERFPERANIGFMQ VVKREHIRLRVYERGAGETQA CGSGACAAVAVGIQQGLLAEE VRVELPGGRLDIAWKGEELQET LTELDDRAVVDYLIKNPEFFIRN ARAVEAIRVPHPVRGTVSLVE WHMARARNHIHVLEENMALL MEQAIANEGLFYRLLYLQRSLT AASSLDDMLMRFHRWARDLG LAVAMPVTINKGNQIIENQLDT ETITISYEFILCQHLIANTEFSYL ALPENYNRLCLPNSKNQTNNRF KTLNSKAIGRLLAAGGVYNGNI EGFRDTAEKLAVPPVQLLTGW CVTAKALPDISAISALTAVKHG NCSSLYTASOPAGO

SEQ ID NO:	SEQ ID NO: of peptide sequence	ŧ .	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22517	52885	A	22645	1479	1793	TRYRSCPSCSATDGVVRNGKST AGHQRYLCSHCRKTWQLQFTY TASQPAGQ*RRNACQYRTPAV VASRQKRWHGDLGMIGSRDV MLFISDSVGAKELDLIIPRLEDK SIALLAMTGKPTSPLGLAAKAV LDISVEREACPMHLAPTSSTVN TLMMGDALAMAVMQARGFNE EDFARSHPAGALGARLLNKVH HLMRRDDAIPQVALTASVMDA MLELSRTGLGLVAVCDAQQQV QGVFTDGDLRRWLVGGGALTT PVNEAMTVGGTTLQSQSRAIDA KEILMKRKITAAPVVDENGKLT GAIKLQDFYQGRGLFNPSIPRVS TADAGLANRARRNWVGSSWK QVGQIILQGGPQNAEKSGQRYT VSHDRHRDAERQRVKSGLPRY AECSTYPSCPVALDLAFASHGI HMHAPGEALRMLVTAMDKAA DARTKLARLLATKGITHEIQIPD ISTKEKAQQPIGLNMEQIKPEKQ DFIKPVIPQAGKHNGVNMSVLR SLLTAGVLASGLLWSLNGITAT PAAQASDDRYEVTQQRNPDAA CLDCHKPDTEAVPPVQLLTGW
						CVTAKAPPDISAISALTAVKHG NCSSLTPLLNPPGSDVIVCAELD EQWGYVGAKSRQRWLFYAYA QSPRTTRCCARIRLHALWRRW

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SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22518	52886	A	22646	807	1350	NRGCPSCSATDGVVRNGKSTA GHQRYLCSHCRKTWQLQFTYT ASQPAGQ*RLLNRLIEPTRGQV LIDGVDIAKISDAELREVRRKKI AMVFQSFALMPHMTVLDNTAF GMELAGINAEERREKALDALR QVGLENYAHSYPDELSGGMRQ RVGLARAINPDILLMDEAFSAL DPLIRTEMQDELVKLQAKHQR TIVFISHDLDEAMRIGDRIAIMQ NGEVVQVGTPDEILNNPANDY VRTFFRGVDISQVFSAKDIARRT PNGLIRKTPGFGPRSALKLLQD EDREYGYVIERGNKFVGAVSID SLKTAAVPPVQLLTGWCVTAK APPDISAISALTAVKHGNCSSLT PLLNPPGSDVIVCAEMDEQWG YVGAKSRQRWLFYAYDSLRKT VVAHVFGERTMATLGRLMSLL SPFDVVIWMTDGWPLYESRLK GKLHVISKRYTQRIERHNLNLR QHLARLGRKSLSFSKSVELHDK VIGHYLNIKHYQ
22519	52887	A	22647		1270	MTAATEQLNASRGLSTREMER QAELAKITTDYINSGSEGDEK LQNMIKAQNDYYAAEDAKRA DWLAGAESAFADYGDSAMDM YGNVNEIASSALNGMTDMMV QFLTTGKANFADFAKNIISMIIK MIAQMVIFNTSLRNDGRSVSSG GAAVSLAAAAGSVATSGFNAS NSAPKVVTIREVERSVDVSGME ASSTSWITIRSFTAETSTFGPTM TASMTFLPTYCSIIASLLSSFKLA AATDGVVRNGKSTAGHQRYLC SHCRKTWQLQFTYTASQPALA RNPNRDITIYWGGREEQHLYDL CELEALSLKHPGLQVVPVVEQP EAGWRGRTGTVLTAVLQDHAT DGVVRNGKSTAGHQRYLCSHC RKTWQLQFTYTASQPGTHQKII DMAMNGVGCRATAR\IMGVGL NTIFRHLKNSGRSR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22520	52888	A	22648	381	945	GAPVASVSISCPSCSATDGVVR NGKSTAGHQRYLCSHCRKTWQ LQFTYTASQPAGQ*RTENQTPH VLTHKWELNYENPWTQGGEHL TMGLVVGWRGLQEGDPARTN LSYYQANVGSGQHRENKGPIN
						LGNVGKSNGERGGKSLYRCKIF LESSVTMKDRELPPVYDGVFEV LQWLLFLSAVPPVQLLTGWCV TAKAPPDISAISALTAVKHGNC
						SSLTPLLNPPGSDVIVCAEMDE QWGYVGAKSRQRWLFYAYDS LRKTVVAHVFGERTMATLGRL MSLLSPFDVVIWMTDGWPLYE
22521	52000	Α.	22640	254	1110	SRLKGKLHVISKRYTQRIERHN LNLRQHLARLGRKSLSFSKSVE LHDKVIGHYLNIKHYQ
22521	52889	A	22649	354	1119	RRWRRGAHEAATGGTGRGPAE EAAPGEPAAGRAAPKSGGKRL GGVLGRGGVGRRSGSPDRSGA HSAAPRRCRQLPGRCPSCSATD GVVRNGKSTAGHQRYLCSHCR KTWQLQFTYTASQPAGQ*RSK GAIGEGPPDPRGIVEQNLGGDIS IPCYNGPRPRRSAQKTTLGPLN QPRIPSAQCAGAFPRVAAAGAG GGGGGGSGGSGFCLGWRLRRR RQRRRWRRGEGEGDGDAGH MRPRPAGRAEARRRRLRGNP PPGARLPNREGNGSGVSWAAE VSGDGRGARTEAARTRRPGG ADNYPGAVPPVQLLTGWCVTA KAPPDISAISALTAVKHGNCSSL TPLLNPPGSDVIVCAEMDEQW GYVGAKSRQRWLFYAYDSLRK TVVAHVFGERTMATLGRLMSL LSPFDVVIWMTDGWPLYESRL KGKLHVISKRYTQRIERHNLNL RQHLARLGRKSLSFSKSVELHD
22522	52890	A	22650	23	658	KVIGHYLNIKHYQ VISRPGSSIQVRGLLLSRHGLAL GSAPDLWKLPWVNLVQGMGQ QSQGSSCREGCSRPQCPSAAPC GAAPAHFCSAATFWESEVQSIP R/CRATDGVVRNGKS/TAGHQR YLCSHCRKTWQ/LQFTYTASQP EVIRSSPTSAPRSDGTKTVPPLM GISP*SAHHRLATSDPSLCLSES LCSREPAAVPRASNQQPQGVK DVHHQFTARTSRYPSKC
22523	52891	A	22651	926	1093	DAULING INKLOVILLOVC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	•	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				Sequence		
22524	52892	A	22652	343	817	GGARSRQNPRCGPCMVVVPGQ SGRSVSADSSTEFREQGGS\AI\S FPKSINRAGPELFRFMENPFNSV *YRKNVRELTPAIMAVLPVEFR TRLAPQNDTMSLIASAMKECSE AKQAVLLNAPEHQKMKEVSEG IASLFRLMPEQVGPLMTMVTS MLGVI
22525	52893	Α	22653	115	425	
22526	52894	Α	22654	747	1173	
22527	52895	Α	22655	1	2709	
22528	52896	Α	22656	1	2583	
22529	52897	A	22657	636	1029	APSDQKRDFLVPHGADSAMAK SG*CESGNDLLSSSHMNQLR/A/ ESIPEDVIAGASALVLTFIIKFVI AENPQWWQQFLKDHVSILAMN EDEAEALTGESDPLLASDKALD WVDLVLCTAGPIGLYMAGGGD VN
						AGVAARSGRNDGTGRCGLCHS CSRYGDHHRQSDRIYRHYDAG RSPSGAVDYGTQRGNRFSRAV YPVGA/RRWR*GLPLV/QQFQPE NETSAAW\VVGINQPLVDIEAK VDDEFIERYGLSAGHSLVIEDD VAEALYQDVQQKNLITHQSAG GTIGNTMHNYSVLADDRSVLL GVMCSNIEIGSYAYRYLCNTSS RTDLNYLQGVDGPIGRCFTLIG ESGERTFAISPGHMNQLRAESIP EDVIAGASALVLTSYLVRCKPG EPMPEATMKAIEYAKKYNVPV VLTLGTKFVIAKNPQWWQQFL KDHVSILAMNEDEAEALTGESD PLLASDKALDWVDLVLCTAGPI GLYMAGFTEDEAKRKTQHPLL PGAIAEFNQYEFSRAMRHKDC QNPLRVYSHIAPYMGGPEKCNI LGRSGWLLYGRGLDKQRLTQY QSKLGAAMVIVAAWCVEDYQ VIRLAGSLTARATRLAHEAQLD VAPLGKIPHLRTPGLLVMDMDS TAIQIECIDEIAKLAGTGEMVAE
22531	52899	A	22659	430	586	VTRTGDARRTRFYRQPAQPCG
22532	52900	A	22660	1	131	
_2002	152550	ı. <u>. </u>	1-2000	<u>ı. </u>	L:	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22533	52901	A	22661	li li	1282	MLPAHLDACVTTLQKIRWTLS
						VVDPTLALRVPLPALSIRDDPPL
						RAYLPGRGALKVIRFDDMDFA
						TIDGINRMLVDIDTNYLFLPRVT
						SSTIRCLSVPTRRTVPAVTASGR
						SVKYYPPDFDPSKIPKLKLPKD
	•					RQYVVRLMAPFNMRCKTCGEY
						IYKGKKF\NARKETVQNEVYLG
						\LPI\FRFYIK\CTRCLGRDSPFKT
						DP\ENTDYTM/ESLGATRNFQG*
						EASWEEEKRV\QK\EREDE/ES
						LNNPMKVLENRTKDSKLEMEV
						LENLQELKDLNQRQAHVDFEG\
						MLRQHRLSEEERRSQQQEEFEQ
						KTAALLEEARKRRLLEDSDSED
						EAAPSPLQPALRPNPTATLDEA
	•					PKPKRKVEVWEQSVGSLGSRPP
			į			LSRLVVVKKAKADPDCSNGQP
		İ				-
						QAAPHPRSPAEQEGGQPYTPDA
22534	52902	_	22662	1	412	WRVLPEPTGCIPGQ
22535	52902	A A	22663	298	532	MGSWSLPGRTLWKEKVRFLKS
22333	32903	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	22003	290	332	KERTNPGSNFFSFPGSQHTREER
		ĺ		1		SQECLSLPVLPSYLATRLSLPDS
		l				CPLKGRKAH*QRHSEI*LMGFE
		1				PGWPGKL*GWIELAFLLGCLHL
		ŀ				TLPLPCPTPLPPKSTKSGCFSLSI
		l				ALCFNKQFAVTK
22536	52904	A	22664	1	193	ALCFNKQFAVIK
22537	52905	В	22665	55	281	
22538	52906	A	22666	86	159	
22539	52907	A	22667	63	1036	VRGGLAAGRGRGSAGAAPVVV
22337	32701	 ^`	22007	03	1030	AAMLGAWAVEGTAVALLRLL
		1				LLLLPPAIRGPGLGVAGVAGAA
						GAGLPESVIWAVNAGGEAHVD
į						VHGIHFRKCIPL*GRVGRALYY
		ł				GMKLPILRSNPEDQSCYQTE\RS
		1				N\EETFGYESPIQREGGLTCWSL
		l				KFAE\VYFAQSQQKVFDVRLNG
		•				` ` `
		ł				HVVVKDLDIFDRVGHSTAHDEI
		1]	IPMSIRKGKLSVQGEV\ST\FTQG
						KLLPLSFSKGYL*PIPKVCAL\YI
						MAG\TVDDVPKASAFNPGLEK
						KEEEEEEEYDEGSNLKKQTNK
						NRVQS\GPRTPNPYALDNSSLM
	50000		22.652	1.570	1550	FPILV\AFGVFIPTLFCLCRL
22540	52908	A	22668	1570	1773	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22541	52909	A	22669	62	454	NKQLNKKRRRTMIALIQRVTRA
						S/VTVEGEVTGEIGAGLLV/LLG
						VEKDDDEQKAN/RDAEGQM/N
		1				LNVQQAGGSVLVVSQF/TLAAD
						TERGMRPSFSQGHHRIRAQAFY
						DYFVES/CRQQR*YAQGGLSAE
						FRLLNS
22542	52910	Α	22670	1	1930	MAGKPEYDKTISTSIVLNALNA
						LGVSAEASGRNDLVVKTVEGD
						RKVSGSAYRETKDRGFHHGTL
						LLNADLSRLANYLNPDKKKLA
]	AKGITSVRSRVTNLTELLPGITH
						EQVCEAITEAFFAHYGERVEAE
						IISPNKTPDLPNFAETFARQSSW
						EWNFGQAPAFSHLLDERFTWG
				-		GVELHFDVEKGHITRAQVFTDS
İ		1		,		LNPAPLEALAGRLQGCLFSLYN
					•	SDLNTVIDNVLRIFPLLLSWISF
		1				WLLYSIVPTIRVPNRDAIVGAFV
						AALLFEAGKKGFALYITMFPSY
						QLIYGVLAVIPILFVWVYWTCV
				1		TVEGEVTGEIGAGLLVLLGVEK
						DDDEQKANRLCERVLGYRIFSD
						AEG\KMNLNVQQAGGSVLVVS
						QFTLAADTERGMRPSFSKGVID
						1 ~
ŀ	1					DIGFNTEISQTRFHQASHLFKRK
						GIYRLHFARRRCQQIQRRQRRR
						RFMQRTAVAITHQQRMKQGIL RLWRNWLLWLLILLLRRNRLL
ł						RLLILLWRNRLLRLLILWLRR
						NRLLWLLILRLRRNWLLRLLIL
						LLRCNRLLRLLYRFNNWRKQR
		ŀ				FVWTIIQQRFHWRKFKIIQRERI
		ĺ				RPSRLFAGITLLLSLRKLTSQFS
						LCRSICIFCVYGHIMLILFAIALL
00540	52011	 	22671	277	462	LCLGELTCQFILLHRIDFIIVVLI
22543	52911	Α	22671	377	463	QTISQTTVQSN*NSGLRISPKTA
22544	52912	A	22672	129	615	QLHGD KARQKSCQPVTTDTQCRHTSSS
22344	34714		22012	149	013	IGKRLAFCONLTOIHTESTMKP
Ī		1				WILLLVMFISGVVMLLPVLGSF
					1	WNKDPFLDMIRETEQCWVQPP
1]			YKYCEKRCTKIMTCVRPNHTC
						i
1						CWTYCGNICLD/IRRAP*/HQC*
						TPRFYWPITTVG*RMTIWSKKS
22545	52012		22673	2	412	AAYFLLGTPTWL
22545 22546	52913 52914	A A	22674	3	390	
22340	134314	ΙΛ.	122014	<u> </u> 4	230	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22547	52915	Α	22675	180	507	CCGGRARSELCLLNLCIKGAD*I
		ŀ				QAW*KFCQWPCERADGRGCRK
		١.				AER\CPCPKGAYSL/YCRESVCC
						LPSFVNKNSSSLCVAHNWGRH
						CYLWTAADVKQFISSHMFPVE
						HL
22548	52916	Α	22676	3	227	GGGRARSELCLLNLCIKGAD*I
						QAW*KFCQWPCERADGRGS*F
						DRGAEVCNQTDWQSIRSSQGG
		ŀ				RDQRFSTERA
22549	52917	Α	22677	567	842	
22550	52918	Α	22678	1	1400	MGWGQGPDVQLMNGHERSPS
						HSPREAKKSGSPLPTDIHCFSPQ
						SQPAMVITGVPTAPAGASAKM
						GLCATPITGACPLWLRGFRGWR
ł						CEDRCEQGTYGNDCHQRCQCQ
1				1		NGATCDHVTGECRCPPGYTGA
						FCEDLCPPGKHGPQCEQRCPCQ
						NGGVCHHVTGECSCPSGWMLS
						FPGWRPI*FSKSL*MQGTVCGQ
						PCPEGRFGKNCSQECQCHNGGT
						CDAATGQCHCSPGYTGERAAV
						PDVRK\CQDECPVGTYGVLCAE
						TCQCVNGGKCYHVSGACLCEA
						GFAGERCEARLCPEGLYGIKCD
						KRCPCHLENTHRGDDREKAEF
						GHLLIALEVQYTEISNCIEKSQE
						ASCYDIRMLKQPYGETNTEEEL
						AVNYVYSIYASKEQIRYKHGRS
						FANGHVKELMEGPSRRIAGSGG
						EKYKRDEEKNTLLRGQNRGGR
						GKNPKKTKAWVKIQGQNKIEIS
						GVGADRKRFFREK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22551	52919	A	22679		2210	MEQKGEFKNDTLIYEFFHIVAK PHPLPIKSSTYTTFQSARVTRFF LDAEKECRCQEGQVEAALIAM GGTCQGTQRLSTAGESESATH MREPAVQCRRDSQCVSVLAGR RRFFQGAASPLHPASVALTQLH VRWASTATIGQVSGFPGKIRVS PTTNLGDSDYIFTASHWGLGHY GPAVHVHSQGGIDERGSNDSG RLYVTNPCGNVPSWTPNIDRLA EDGVKLTQHISAASLCTPSRAA FLTGRYPVRSGMVSSIGYRVLQ WTGASGGLPTNETTFAKILKEK GYATGLIGKWHLGLNCESASD HCHHPLHHGFDHFYGMPFSLM GDCARWELSEKRVNLEQKLNF LFQVLALVALTLVAGKLTHLIP VSWMPVIWSALSAVLLLASSYF VGALIVHADCFLMRNHTITEQP MCFQRTTPLILQEVASFLKRNK
						HGPFLLFVSFLHVHIPLITMENF LGKSLHGLYGDNVEEMDWMV GRILDTLDVEGLSNSTLIYFTSD HGGSLENQLGNTQYGGWNGIY KGGKGMGGWEGGIRVPGIFRW PGVLPAGRVIGEPTSLMDVFPT VVRLAGGEVPQDRVIDGQDLL PLLLGTAQHSDHEFLMH\YCER VLHAARWHQRDRGTMWKVHF VTPVFQPEGAGACYGRKVCPCF GEKVVHHDPPLLFDLSRDPSET HILTPASEPVFYQVMERVQQAV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
·	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
22552	52920	Α	22680	185	2091	GSTERDNMLHLHHSCVCFRSW
						LPAMLAVLLSLAPSASSDISASR
						PNILLLMA\DDHGVGDIGCYGN
		l				NTMRTPNIDRLAEDGVKLTQHI
İ		ĺ				SAASLCTPSRAAFLTGRYPVRS
						GMVSSIGYRVLQWTGASAGFT
						TN\ETTFAKILKEKGYATGLIGK
		ł				WHLGLNCESASDHCHHPLHHG
						FEHFYGMPFSLMGDCARWELS
						EKRVNLEQKLNFLFQVLALVA
		l	•			LTLVAGKLTHLIPVSWMPVIWS
						ALSAVLLLASSYFVGALIVHAD
						CFLMRNHTITEQPMCFQRTTPLI
						LQEVASFLKRNKHGPFLLFVSF
						LHVHIPLITMENFLGKSLHGLY
						GDNVEEMDWMVGRILDTLDV
						EGLSNSTLIYFTSDHGGSLENQL
						GNTQYGGWNGIYKGGKGMGG
						WEGGIRVPGIFRWPGVLPAGRV
						IGEPTSLMDVFPTVVRLAGGEV
:						PQDRVIDGQDLLPLLLGTAQHS
						DHEFLMHYCERFL\HAARWHQ
						RDRGTMWESPTL*PPCVSSQEG
						AGAC\YGRK\VCPCFGEK\VVHH
						DP\PLL\FDLSRDPSETHILTPASE
						PVFY\QVMER\VQQAVWEHQR
						TLSPVPMQLDRLGNIWRPWMQ
						PCCGPFPLCWCLREDDPQINVC
						SEKLEPRFLNFVTQIETNQLAM
22553	52921	A	22681	209	392	
22554	52922	A	22682	1	513	
22555	52923	A	22683	3	613	
22556	52924	Α	22684	1	2115	

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SEQ ID NO:	SEQ ID NO: of peptide sequence	ł .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22557	52925	A	22685	53	1966	VGQPRGLRTLEKLTASVAGGA ANFFGDIERDLGPRDTSRGIRG GNLGDRGCSSAEMAYASWQR WSPVEWARWMWTAVTSSGDS SLLVLQGDSGKRSSDSEEAFET PESTTPVKAPPAPPPPPPEVIPEP EVSTQPPPEEPGWGSETVPVPD GPRSDSVEGSPFRTQAHSFSGVF DEDQPIASTGTDILDFDNIELVD TFQTLEPRASDAKNQEGKVNT RRKSTDSVPISKSTLSRSLSLQA SDFDGASSSGNPEAVALAPDAY STESAKRVVPPASGGGRVQNSP PVGRKTLPLTTAPEAGEVTPSD SGGQEDSPAKGLSVRLEFDYSE DKSSWDNQQENPPPTKKIGKKP VAKMPLRRPKMKKTPEKLDNT PASPPRSPAEPNDIPIAKGTYTF DIDKWDDPNFNPFSSTSKMQES PKLPQQSYNFDPDTCDESVDPF KTSSKTPSSPSKSPASFEIPASA MEANAV\DGDGLNKPAKKKKT PLKTMVEDVMSVCSLFDTFRV KKSPKRSPLSDPPSQDPTPAATP ETPPVISAVVHATDEEKLAVTN QKWTCMTVDLEADKQDYPQPS
						DLSTFVNETKFSSPTEELDYRNF YEIEYMEKIGSSLPQDDDAPKK QALYLMFDTSQESPVKSSTVR

	odon for last amino acid *=Stop codon, /=possible nucleotide
sequence 09/340,217 codon for peptide of	f peptide sequence deletion, \=possible nucleotide insertion)
Sequence	
22558 52926 A 22686 662 29	927 HLRRKRNLPRREGLSPALLEET
	PLEPAVGPKAACPLDSESAEGV
	VPPASGGKRCLFTTAPEAGEVT
	PSDSGGQEDSPAKGLSVRLEFD
	YSEDKSSWDNQQENPPPTKKIG
	KKPVAKMPLRRPKMKKTPEKL
	DNTPASPPRSPAEPNDIPIAKGT
	YTFDIDKWDDPNFNPFSSTSKM
	QESPKLPQQSYNFDPDTCDESV
	DPFKTSSKTPSSPSKSPASFEIPA
	SAMEANGVDGDGLNKPAKKK
	KTPLKTDTFRVKKSPKRSPLSD
	PPSQDPTPAATPETPPVISAVVH
	ATDEEKLAVTNQKWTCMTVD
	LEADKQDYPQPSDLSTFVNETK
	FSSPTEELDYRNSYEIEYMEKIG
	SSLPQDDDAPKKQALYLMFDT
	SQESPVKSSPVRMSESPTPCSGS
	SFEETGSPLRNTAA\KNQ\HPVP
	TRDWAP*PRSSHLAGCQRKSSQ
	E/VELGGPWGLGHPFQEGDLKF
	TASPRGSFASADALLSRLAHPV
	FSL/CGALDYL\EPNFRQKRNPP
	LFAQKLQREA\VHPTDVSI\SKT
	ALYSRIRTTEVEKPAGLLFQQP
	DL\DSALQIARAEII\TKEREVSE
	WKDKYEESR\REVMEMRKI\LA
	EYEKTIAQMIEDEQREKSVS\HQ
	TVQQLVLEKEQA\LADLNSVEK
	\SLADLFRRYEKMKEVLESFRK
	NEEVLKRCAQEYLSRVKKEEQ
	RYQA\LKVHA\EEKLDRANAE\I
22559 52927 A 22687 374 12	268 AETNMVWYDWMRPSHAQLHS
	DYMQPLTEAKAKSKNKVRGV
	QQLIQRLRLIKSPAEIERMQIAG
	KLTSQAFIETMFTSKAPVEEAFL
	YAKFEFECRARGADILAYPPVV
	AGGNRSNTLHYVKNNQLIKDG
	EMVLLDGGCESSCYVSDITRTW
	PVNGRFTAPQAELYEAVLEIQR
	DCLALCFPGTSLENIYSMMLTLI
	GQKLKDLGIMKNTQGK*CL/RR
	LLE/QYCPHHVGHYLGMDVHD
	TPDMPRSLPLQPGMVITIEPGIYI
	PEDDKDAPEKFRGLGVRIEDDV
	VVTQDSPLILSADCP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22560	52928	IA	22688	329	1389	RIWKTQMMRMKKMMMMMM
						MMMRTTLFFQQESQEAIFFPFD
·		l]		LFPMCPFGCQCYSRVVHCSDLG
ŀ						LTSVPTNIPFDTRMLDLQNNKI
						KEIKENDFKGLTSLYGLILNNN
ŀ						KLTKIHPKAFLTTKKLRRLYLS
						HNQLSEIPLNLPKSLAELRIHEN
						KVKKIQKDTFKGMNALHVLEM
						SANPL**/SMGIEPGAF*RG*RCS
	1					/YIRIAEAKLTSV\PKGL\PPTLLE
		1			,	LHLDYNKISTVELEDFKRYKEL
						QRLGLGNNKITDIENGSLANIPR
						VR\EIHLGNIKLKKIPSGLPELKY
						LQIIFLHSNSIARVGVNDFCPTV
}						PKMKKSLYSAISLFNNPVKYWE
<u> </u>	ļ	ļ			1.500	MQPATFRCVLSRMSVQLGNFG
22561	52929	A	22689	1	4608	
22562	52930	A	22690	1	465	
22563	52931	A	22691	1	1542	EAVI CSCRCRRRSCARDCCRNA
22564	52932	A	22692	1	4226	EAVLCSSRGRPDSSMPDCCRNA
						ALEARSCQSMTSLFSNTVSPTQ
						DGTSSLPRRQSSFAKPPLRALY
			ŀ			DLLIAPMEGGVVESTEPLGTSR
						SGFASGTALFWLGDNGKAASP
		ŀ				PGVSVSSLENEDDNTQCNRLEQ
						GYREAAVPDWTQDLALCLLPT
						LQSLKARTAPHLCVPSSCYKA
						WYTLAVSKVSGMVEGKAWIIL
						LLLAASTHALLQRPQEGDITPL
		1				AGSRAAAVSCQARPFSTFSGLM
	<u> </u>					HSSGPVGRHRQLILVLE
22565	52933	Α	22693	2	4197	ALDFPGRRFRPTASFIWASVLFE
	1					TIRHEAEVSTDYKLSLFDLQTSS
		1				YQALQRVLVSLGHHDEALAVA
	1		}			ERGRTRAFADLLVERQTGQQD
			}		:	SDPYSPVTIDQILEMVNGQRGL
						VLYYSLAAGYLYSWLLAPGAG
		1				IVKFHEHYLGENTVENSSDFQA
		1				SSSVTLPTATGSALEQHIASVRE
		1				ALGVESHYSRACASSETESEAG
		1				DIMDQQFEEMNNKLNSVTDPT
						GFLRMVRRNNLFNRSCQSMTS
			1			LFSNTVSPTQDGTSS
22566	52934	A	22694	1	618	
22567	52935	Α	22695	1	3528	
—— —						L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22568	52936	A	22696	85	1193	RPIEGQLGTCHPGSEPRTIRWRA PGKGLRLYTHPPTSDIAGRTGA AASFPAPLHLPREPRAGVDGTD AERILYPRRAAPAASGVGSRPA GEAVRADEAGAAGLGGFQSER NRSWGGESLGSRQPQKMPAFN RLFPLASLVLIY*VSVCFPVWGE \VPLENGGRARANPMKLRCISC MKREEVEATTVVEWFYRPEGG KDFLIYEYRNGHQEVESPFQGR LQWNGSKDLQDVSITVLNVTL NDSGLYTCNVSREFEFEAHRPF VKTTRLIPLRVTEEAGEDFTSV VSEIMMYILLVFLTLWLLIEMIY CYRQVSKAEEAAQENAPGNSL EGGSQGLQINSTQDGLVTGQQ
22569	52937	A	22697	277	527	DVGSHRAREHMPDFEEPPSAK VRVSWEYFCAPGPAGSPAGPG RGEEKQRLGLLIRDARSPAPPG PPVGSVRG/PKPAPAGRVRAKG RRA*GAACSLAAGAWVEPGI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22570	52938	Á	22698	226	2268	SVKNYTKCHVRNEQICNKLTSC
						KSCSLNLNCQWDQRQQECQAL
						PAHLCGEGWSHIGDACLRVNSS
						RENYDNAKLYCYNLSGNLASL
						TTSKEVEFVLDEIQKYTQQKVS
İ						PWVGLRKINISYWGWEDMSPF
						TNTTLQWLPGEPNDSGFCAYLE
						RAAVAGLKANPCTSMANGLVC
						EKPVVSPNQNARPCKKPCSLRT
						SCSNCTSNGMECMWCSSTKRC
						VDSNAYIISFPYGQCLEWQTAT
						CSPQNCSGLRTCGQCLEQPGCG
						WCNDPSNTGRGHCIEGSSRGPM
						KLIGMHHNEMVLDTNLCPKEK
						NYEWSFIQCPACQCNGHSTCIN
						NNVCEQCKNLTTGKQCQDCMP
						GYYGDPTNGGQCTACTCSGHA
						NICHLHTGKCFCTTKGIKGDQC
					·	QLCDSENRYVGNPLRGTCYYK
					İ	T\LIDYQFTF\SLLQEDDRHHTAI
						YFIVNPKQSNKNLHISINASNNF
						NLNITWPC*STAGTISGEETYIV
						SKNNILEYRDSFSYEKFNFRSNP
						NITFYVYVSNFSWPIKIQIAFSQ
ļ						HNTIMDLVQFFVTFFSCFLSLLL
						EAAVVWKIKQTCWASRRREQL
						LRERQQMASRPFASVDVALEV
						GAEQTEFLRGPLEGAPKPIAIEP
						CAGNRAAVLTVFLCLPRGSSGA
İ						PPPGQSGLAIASALIDISQQKAS
						DSKDKTSGVRNRKHLSTRQGT
22571	52939	В	22699	252	360	25125K156VKKWARIESTKQ61
22572	52940	A	22700	1	84	
22573	52941	A	22701	1	404	
22574	52942	A	22702	86	292	NPEMRCREFCLPSCRPELALE*
22571	327 12	l' .	22,02			G*SVRDVA*AQMCSVVWAGRP
		ŀ				LGFAA*S*GSGHSCGRCSIWGE
						CWG
22575	52943	A	22703	82	379	HTLQPFISMGSLTKMFKVISEGP
						MD*RPHPKVRPMT*HSTPAPAN
1						FMGHFFWKIMICTDRTEVATLV
						AGVVSGASVGGKSPPYRKWNS
]						PEPDVKGPRVL
22576	52944	A	22704	2	382	
	1-27.1	1 .	L, 0 T	<u> </u>	1002	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22577	52945	Α	22705	1	479	HMPSGETAPLRIVDTGTGTGTV
		ł				\AYAPTE\LGLHEMHIKYMASHI
						LESPLRFYVTTPTVPSVSAYGPG
				,		LVYGVAN/NTATFTIVTEDAGE
						GGLDLAIEGPSKAEEPRPRLSRA
						PGVDGAGASPGLGEDTGPQAP
						GLQVLTERIPGPSAPTPEFLSVG VSAVC
22578	52946	Α	22706	1	590	
22579	52947	A	22707	171	406	
22580	52948	A	22708	269	2477	
22581	52949	В	22709	167	7472	
22582	52950	Α	22710	l	520	
22583	52951	A	22711	2218	2472	
22584	52952	Α	22712	722	1555	SRNWYEALNNTWGREDICVVD
		ĺ				GWQSFRRQDYAFLQERGSGGD
						LWSSTLAQAMTLVESRHPRHQ
					,	ALSTQLLDAIMPYCGNTLRLGV
						TGTPGAGKSTFLEAFGMLLIRE
						GLKVAVIAVDPSSPVTGGSILG
						DKTRMNDLARAEAAFIRPVPSS
						GHLGGASQRARELMLLCEAAG
						YDVVIVETVGVGQ\SETEVARM
						VDCFISLQIAGGADDLQGIKKG
		l				LMEVADLIVINKDDGDNHTNV
						AIARHM/YESALHILRRPPMHR
				1		VPRNPKVNLLPGTAMANLLS
22585	52953	В	22713	381	425	
22586	52954	Α	22714	1	1185	
22587	52955	A	22715	1	3009	
22588	52956	Α	22716	22	527	YEGIVMKPLIIVNPADCIGC\RT
						CE\VACVVAHPSEQELNADVFL
		1				PRLKVQRLDSISAPVMCHQCEN
		l				APCVGACPVGALTMGEQVVQT
					ļ	NSARCIGCQSCVSACPFGMITIQ
						SLPGDTRQQIVKCDLCEQR*EG
				1		PACVESCPTQALQLLIERELRRV
						RQQRIVVTGENPL
22589	52957	Α	22717	187	325	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22590	52958	A	22718		2240	MRSDPDTSVNEIETKLSALLGS ETTGEILFDLLCANGPEWNRFV TLEMKYGRIMLDTAKIIDEQDV PTHILSKLTFTLRNHPEGVVMK NFEVLQPLQNSLSGLPLWVSER ILQQINQLTHYEPVIGIMGKTGA GKSSLCNALFAGEVSPVSDVAA CTRDPLRFRLQIGEHFMTIVDLP GVGESGVRDTEYAALYREQLP RLDLILWLIKADDRALATDEHF YRQVIGEAYRHKMLFVISQSDK AEPTSGGNILSTEQKQNISRKIC LLHELFQPVHPVCAVSVRLHPV VALLQGFRTDDERTHRYHYDS QHRLVDYTRTQYEEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRSRIQTIYQPGSFTPLIR VETATGELAKTQRRSLADALQ QSGGEDGGSVVFPPVLVQMLD RLESEILADRVSEESRRWLASC GLTVEQMQNQMDP\TEGTTAW Y/AEYDEWGNLLNEENPHQLQ QLIRLPGQQYDEESGLYYNRHR YYDPLQGRYITQDPIGLKGGW
:						NFYQYPLNPISDIDPLGLSMWE VARQMNAIASKLPEGFHIGINFS ASHIISPTFVDECLNFRDSFTRR DLNLVLEVTERDLLNVDESLVQ RLNILHENGFVIALDDFGTGYS GLSYLHDLHIDYIKIDHSFVGRV NADPESTRILDCVLDLARKLSIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
					loos:	
22591	52959	Α	22719		2301	MALTSGVHTGEKPYKCAQCEE
						AFSRKSELIIHQIIHTGEKPYECT
						ECGKTFSRKSQLIIHQRTHTGEK
						PYKCTKCGKSFCQQSHLIGHQR
	1	•				IHTGEQPYLGLSVAERPKEYED
						DRALKMAAPCGNSSREQPCCFL
ļ						NKASEDAERAKPPS*EPGRLRT
ļ						P*VAQPH*VPHFMATWVRKHP
	:					VHDRQFRLHGQIRELNWDVVR
						PEQFTGVATNCSGITYPQGGWL
						CPAELTRNVLELAQQQGLQIYY
						QYQLQNLSRKDDCWLLNFAGD
ł						QQATHSVVVLANGHQISRFSQT
						STLPVYSVAGQVSHIPTTPELAE
	İ				:	LKQVLCYDGYLTPQNPANQHH
			i			CIGASYHRGSEDTAYSEDDQQQ
						NRQRLIDCFPQAQWAKEVDVS
	ŀ					1 -
						DKEARCGVRCATRDHLPMVGN
	İ					VPDYEATLVEYASLAEQKDEA
						VSAPVFDDLFMFAALGSRGLCS
						APLCAEILAAQMSDEPIPMDAS
						TLAALNPNRLWVRKLLKGKAV
						KAG
22592	52960	Α	22720	1021	1734	
22593	52961	Α	22721	1247	2412	RRSHHAANLDIRIRMPQTRQVT
			ł			LPDPELHPDSTLSMWPD/NRIAR
						DAHYLYRYDRHGRLTEKTDLIP
						EGVIRTDDERTHRYHYDSQHRL
		ŀ	İ			VHYTRTQYEEPLVESRYLYDPL
						GRRVAKRVWRRERDLTGWMS
						LSRKPQVTWYGWDGDRLTTIQ
						NDRTRIQTIYQPGSFTPLIRVET
	1					ATGELAKTQRRSLADALQQSG
		ŀ				GEDGGSVVFPPVLVQMLDRLE
		Ì		<u> </u>		SEILADRVSEESRRWLASCGLT
						VEQMQNQMDPVYTPARKIHLY
	1					
				1		HCDHRGLPLALISKEGTTEWCA
				1		EYDEWGNLLNEENPHQLQQLI
		1				RLPGQQYDEESGLYYNRHRYY
		ŀ		1		DPLQGRYITQDPIGLKGGWNFY
		l				QYPLNPVTNTDPLGLEVFPRPFP
						LPIPWPKSPAQQQQWYPTFCC
22594	52962	Α	22722	1	4767	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22595	52963	Α	22723	668	1922	YPHADSPECWLSSSSERGKSSV
İ						TADLSRPRRLRPIPTCHGKPIAC
						TTTKAKNAFRPIAGASATGRLA
						NSPIIILPNAAIRQEFGITLDPEEV
į.						FTFQGVKLYEYRYFPDCVRSRN
1						EAGILVEQILSTLFSVGEHFSLK
						RAPFFKVPSLRSFYAFGGLTTG
						VVRRNGNHGVNTPARKIHLYH
						CDHRGLPLALISKEGTTEWCAE
						YDEWGNLLNEENPHQLQQLIR
1						LPGQQYDEESGLYYNRHRYYD
						PLQGRYITQDPIGLK\GGWNFY
						QYPLNPVQYIDSMGLASKYGH
		İ				LNNGGYGARPNKPPTPDPSKLP
						DIAKQLRLPYPIDQASSAHNVF
						KTFFRALSPYDYTLYCRKWVK
						PNLTCTPQDDSQYPGMDTKTA
						SDYLPQTN/WPVPSSIVSCTTGV
						VP*VVCQWVFAWPPDISHRRTG
22596	52964	Α	22724	25	510	RITDFGVKKIIRVGSCGAVLPHV
						KLRDV/VI/GMGAC/TDSKVNRI
						RFKGQAFGGIADFAMGRTAVD
						PGKALGINARVGNLVSADLFYS
						PDGEMFDVMEKYGILGVEMEA
						AGIYGVAAEFGAKALTICTVSD
						HIRTHEQTTAAERQTTFNDMIKI
						ALESVLLGDKE
22597	52965	Α	22725	415	759	HRRRIVIDDTGIFRPRNTAQQLR
						QRPIAMSASGVVQIILEGHWRT
1						HSLRNGFHRGFRL*SSPKVGMQ
						HGSRQVDHRTQRALRFFRQPGI
						DFLRPVFTPRRQRLAATNHFSR
		<u> </u>				FVEH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	:			sequence		
22598	52966	Α	22726	2	2329	GTLPGCGAMHTCPSGPATHLV
						RRRIVRLPACVVTRANVPTPSR
						PVNATIGAFHSG*MASQGRRLS
						AQYPWSPFQYA\ARPVPTVPGR
						LVFTGTSKAPKISVMGPGMGIP
						SCSISAKELITDFGVKKIIRVGSC
	!					GAVLPHVKLRDVVIGMGACTD
						SKVTRIRFKDHAFAAIADFDMV
						RNAVDAAKALGIDARVGNLFS
						ADLFYSPDGEMFDVMEKYGIL
į						GVEMEAAGIYGVAAEFGAKAL
İ						TICTVSDHIRTHEQTTAAESQTT
	l					FNDMIKIALESVLLGDKEEDRVI
						RFGKARATRYALLRPYRGIERIP
						VWRVDDTGKAHKFADIRLCWP
						QGSCLDVAKAREVIRGSQPLPA
						WCENWAQEVDARLTKEAQSA
						LRPVINLTGTVLHTNLGRALQA
						EAAVEAVAQAMRSPVTLEYDL
						DDAGRGHRDRALAQLLCRITG
						AEDACIVNNNAAAVLLMLAAT
						ASGKEVVVSRGELVEIGGAFRI
	ŀ					PDVMRQAGCTLHEVGTTNRTH
						ANDYRQADFQFDDFVMAFTQN
						INGPTQNTRALHGGHKRPDFLP
						LLGAFHGAIDILAPGGLHGYSL
						PSSFFHALALSAAFPGKRCKLS
						VHLPFWGLKDGGPLLTAPLGG
						ALVGIVGGVGGSNLTFPFHTNF
						LNVRFVIRRISHRHHHFYRIAKL
						LMNGAQGSNMLIAAVAGVPHQ
						NGHFPFTVADRFAIVLSEGSIRV
22599	52967	A	22727	1	702	
22600	52968	Α	22728	876	1318	RCHLANASPAAGSHAPSPDSLS/
1						KIR/QEVG/PPLP/PLLAPLIATPP
-		1				RTSQPLSPLISSSSPSSPASPVGQ
						VSPFRETPVPPAMSPWPEDPRR
						ASPPDPSPSPSAASASERVVPSP
						LQFCAATPKHALPVPGRLPPCA
	l					SGHAMPCWRAKIK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
22601	52969	A	22729	485	1455	PVVQRARNASSCQPFSTSVWAI
						PSISATSVPMCGATHSILSPKKS
						TVSDRIGSMQISFFPLSRSAEK*
						ESPCSSDAFHAIFSVLSGLALHS
						TTTSLCSSTSGQLVCC**TSLPPT
						TYGMIACAAPVE*SPRCPV*PPA
						SAI*RCNNVEALCSTPFERQP*E
	,					PAK\SMPDHKPHECVYVVGE*D
						PAPHPNSREQIHPAHECLAGFP
		1				VTQESLFAPLEIVPAYCCEPGRE
		l		ļ		PRSARSFPFAPRELT*VQLELGT
						GL*RETLDLVVSAGWA*ARRIS
		l				GAVTATPETGTLCQLDSGAECH
		l				YRETGVELP*RSPRPVGHGSSR
	ł	1				HSRRWRT
22602	52970	Α	22730	814	5001	PENSSHRPKPIIPMIASTLINANQ
						NSVSPYKRTFTRLTALMIIKKA
]		<u>.</u>		AAQIHVGTSGNQYWRKLDFNS
		i				PGGSSPVENSDCSTNSRLSFSPE
		i				NILIQNQDIVREAAVQGDGQKQ
		ĺ		1		RQPQATDLDSSGTHGSEMLPAT
						EVTVSGGFSVEETSCGDTGRSG
						GEALAVANDSTSTPQNANGLW
						KLKSTTPGGALPECFGTTDTTFS
		1		}		SAFCRKHGETQDTSQSSLPGTL
						HCYTGIREGGDDTEVESEAFSC
		<u> </u>				SEGSEQQDAPD
22603	52971	A	22731	34	102	
22604	52972	A	22732	584	893	QKLLPNRMGHPVFVKSSQDPV
						LLSRIAWKDASTVTWVIEQDPV
						LRRRRRRTRRRRTRMRRRRGR*
						*/SKRRRMR/GRRGRR*CRRRRR
	<u> </u>					RSSRSRPTRRQRNRNQV
22605	52973	Α	22733	2	269	IGKLYWFMPPSANLALRELGIV
						LFLSVVGLKSGLSWIGYGALIT
				^		AVPLITVGILARMLAKMNYLT
						MCGMLAGSMTDPPA*PRFQTN
						DREEQHDPQLPQREVGAWRHK
						PVQLAD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22606	152074	1	22734	3	958	RCYSLLSRAPHPGIYISIAERV/*I
22606	52974	Α	22734	3	936	
						RLNSANFTLNSLLPAEKKLAEE
						FAVSRMTIRKAIDLLVAWGLV
						VRRHGSGTYLVRKDVLHQTAS
						LTGLGEVLKRQGKTVTSQVLIF
						EIMPAPPAIASQLRIQINEQIYFS
	1					RRVRFV\EGIPLMLEDSYMPVK
ŀ						LFRNLSLQHLEGSKFEYIEQECG
ŀ						ILIGGNYESLTPVLADRLLARQ
						MKVAEHTPLLRITSLSYSESGEF
						LNYSVMFRNATQIQGWRSHFR
						R\LHPKKSLTVPPERAPPVFGGL
						IMRKNHSQRIDSGVRERRRTAG
		i				WMKIISKRQRRRIGHGTCQHPA
	:					HGQVIHFG
22607	52975	Α	22735	2	285	GDEMTTSTLQKAIDLVTKATEE
		l				DKAKNYEEALRLYQHPVEYFL
İ						HA\IKYEAHSDKAKESIRAKCV
			1			QYLDRAEKLKDYLRSKEKHGK
						LCLKDRVT
22608	52976	Α	22736	545	791	
22609	52977	Α	22737	249	1574	NHKAIDLVTKATEEDKAKNYE
						EALRLYQHAVEYFLHAIKYEA
						HSDKAKESIRAKCVQYLYRAE
						KLKDYLRSKEKHGKKPVKENQ
					:	SEGKGSDSDSEGDNPEKKKLQE
						QLMGAVVMEKP\NIRWNDVAG
		1				LEGAKEALKEAVILPIKFPHLFT
						GKRTPWRGI\LLFRPPGT\GNSN
						RPKAGAPKATNSTFFSVSS\SDV
		ŀ				KSKRLGETEKLV*NLF\ELARQ
		1				HK\PSIIFIDEVDSLCGSRNENES\
						QAARRIQNGSSLVQIAGGLGIN
		1				NDGDSGFFGATNIP\WVLD\SAI
						R\RKV*KTKFYIPLPEKACPRPD
						VSRLHLGSTPHN\LRVATFHKL
						ARKTEGYSGADISIIVRDSLMQP
						VRKVQSATHFKKVCGPSRTNPS
						MMIDDLLTPCSPGDPRAMEMT
						WMDVPGDKLLEPVVCMSDML
						RSLATTRPTVNADDLL\KVKKF
						SEDFGQES
22610	52978	A	22738	52	567	
22611	52979	В	22739	1	1293	
22612	52980	A	22740	443	1027	
22613	52981	A	22741	2077	2430	
22614	52982	A	22742	1	504	
	1	ı. •	<u> </u>	<u> </u>	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22615	52983	A	22743	3244	3947	RNEITPFRAMMIIRRMAHRFQT
						EEIKRISFIKRRVGNKDEINPGA
						RWRRFHDDGITFLRLLDAGLTI
						PSGKTIVHPLARQRRGLVKIQFL
						R*LLAKLTRQPLLPLQPVTARL
						CLNPRLRRQRKIRMPAGTKWO
						RLCPLIGDR*RHRQRSAVYALS
						VRHNPVHHILRKNGDMYSFPIT
						IGNNVWIESHVVINPGVTIGDNS
						VIGAGSIVTKDIPPNVVAAGVP
						CRVILEINDRDK
22616	52984	A	22744	742	1158	CKVILLINDRDK
22617	52985	A	22745	439	2882	PDWQAFQSPPFPASLIIHPKPAK
22017	132703	A.	22743	437	2002	LPGWLCIVQWSLLAITRTPTPV
						RWAQQTTETVGLVVGDVSDPF
		,		:		FGAMVKAVEQVAYHTGNFLLI
				•		-
						GNGYHNEQKERQAIEQLIRHRC AALVVHAKMIPDADLASLMKQ
						``
						MPADKQCESIPDQNNQTGTQY
						PRGKEPKLLRRLMHMNRKVGP
						RGPRRMLLRYLDVEFLSVALA
						VVHLGPQGTCSRLFLVARALV
					:	VAVSEMDKSLSAQAQIYFASSN
						GPVPDLAYQVDFPRLEIGL*F\E
						YVDTGAETTLVPGDVLYVPAG
						GWNFPQWQAPATTFSVLFGKQ
						QLGF/SAVVQWDGKQYQNLAK
						QHVARRGPRIGSFLLQTLNEMQ
						MQPQEQQTARLIVASLLNHCR
						DLLGSQIQTASRSQALFEAIRDY
						IDERYASALTRESVAQAFYISPN
						YLSHLFQKTGAIGFNENLSHTR
						LGHA*TLLKGYDLKVKEVAHA
						CGFVDSNYFCRLFRKNTERSPS
						EYRRHRRLHDNRICQFKTPTDN
						PENPORYLTFRRIDKETSSLPKK
						VAPSCKRLSVSRSSPKPSISIRRC
				ļ		MRLMLIEGRRSSFIHPDKDVDG
						FHPYNVGRLCQRAPRLRPCTPR
						GIVTLLERYNIDTFGLNAVVIGA
						SNIVGRPMSMELLLAGCTTTVT
						HRFTKNLRHHVENADLLIVAV
						GKPGFIPGDWIKEGAIVMMSGI
						NRGVATDASCFPLVPFANRVSG
22618	52986	Α	22746	2	126	
22619	52987	Α	22747	1	465	
22620	52988	C	22748	78	308	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22621	52989		22749	<u>l</u> Ii	2797	MELIDDVI AVODVAL DED CEEDE
22021	32969	A	22149	['	2191	MEHPPVLAVQRVALPSRSSEPE
						LFLNPPTDEETEAHRGMYLDQ
						GPRTKDWIPVAPGVREPSPLGP
1						GPSSLWLLVVGGTVEEGSRKH
				1		RPTCKKDFTKSKTVDQAPTLCA
				ł		RHQMLNIQFLISDRDPQCNLHC
						SRTQPKPICASDGRSYESMCEY
						QRAKCRDPTLGVVHRGRCKDA
						GQSKCRLERAQALEQAKKPQE
						AVFVPECGEDGSFTQNTSCIHSS
						VYQIGENSSLTFLEKREESGHL
						QRDWFPLWFPRKARVI
22622	52990	Α	22750	1	5295	MNLNQEEIPDLPEKEFRRLVIKL
						IREGLEKGKAQCKEIQKMIQEV
						KREIFQEIESLKKKPPKIQKTLDI
						LLKMQNALESLSDRTEQAEEIN
						SELEDKVFKLTQSNKDKEKITR
						KYEQSLQEVWSYVKRPNLRIIG
						VPEEEENTKSLENILGGIIEENFP
					ŀ	SFARDLDIQIPEAQRTTGKLIAK
						RSLLRHIVFRLSKVKMKERILR
						AVRQKHQATVSQKNKAGGITL
						PDFKLYYKATVSQKNKAGGITL
						PDFKLYYK
22623	52991	Α	22751	1	891	GSPFEVMPRDDVFFIVDVKI*DI
						GVYSCTAQNSAGSISANATLTV
					:	LETPSFLRPLLDRTVTKGETAV
+			1			LQCIAGGSPPPKLNWTKDDSPL
			ł			VVTERHFFAAGNQLLIIGDSDV
			<u> </u>			SDAGKYTCEMSNTLGTERGNV
						RLSVIPTPTCDSPQMTVL/HLLD
						DDGWGHCGLYVIIAVVCCVVG
						TSLVWVVIIYHTRRRNEDCSITN
						TDETNLPADIPSYLSSQGTLADR
						QDGYVSSESGSHHQFVTSSGAG
						FFLPQHDSSGTCHIDNSSEADVE
		ŀ			•	AATDLFLCPFLGSTGPYVFEGK
		<u> </u>				CVWAQNPF
22624	52992	Α	22752	1	2947	MSAPSLRARAAGLGLLLCAVL
			ĺ			GRAGRSDSGGRGELGQPSGVA
			}			AERPCPTTCRCLGDLLDCSRKR
						LARLPEPLPSWVARLDLSHNRL
						SFIKASSMSHLQSLREVAGNRIV
						EILPEHLKEFQSLETLDLSSNNIS
						ELQTAFPALQLKYLELNRNKIK
]		NVDGLTFQGLGALKSLKMQRN
						GVTKLMDGAFWGLSNMEILQL
						DHNNLTEITKGWLYGLLMLQE
						LHLSQNAINRISPDAWEFCQKL
						SELDLTFNHLSRLDD
22625	52993	A	22753	2	417	
	<u> </u>				·	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22626	52994	Α	22754	1	262	VLDIGS/SLTVPDEFSKQE/RAD
			:			GMWWKQLVAGAVAGAVS/RT
		İ				GTAPLDRLKVFMQIKRAILGQQ
						ETLMCRSASWLAPWLVPQPNH
						HLPYG
22627	52995	Α	22755	1	762	
22628	52996	Α	22756	1	1527	RLAAAGADPAGSRGGRGSREA
	:					PTEAARCRASPPPRAGAMRGSP
						GDAERRQRWGRLFEELDSNKD
	İ					GRVDVHELRQGLARLGGGNPD
						PGAQQGISSEGDADPNGGLDLE
						EFSRYLQEREQRLLLMFHSLDR
	:	İ				NQDGHIDVSEIQQSFRALGISIL
						LEQAEKNFAQA/VDRDGTMTID
						WQEWRDHFLLHSLENVEDVLY
						FWKHSTVLDIGECLTVPDEFSK
						QEKLTGMWWKQLAAGAVAG
						AVSRTGTVPLDRLKVFMQVHA
						SKTNRLNILGGL*SMVLEGGIRS
						LWRGNGINVLKIAPESAIKFMA
						YEQIKRAIPG\E*ETLPVLERFVA
ŀ						GSLAGATAQTII\YPMEVLKTRL
	1					TLRRTGQYKGLAGLRRGGILER
						EGPRAFYRGYLPNVLGIIPYAGI
						DLAVYETLKNWWLQQYSHDS
						ADPGILVLLACGTISSTCGQIAS
						YPLALVRTRMQAQASIEGGPKL
						SMLGLLRHILSQEGMRGLYRGI
						APNFMKVIPAVSISYVVYENMK

2965

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
-	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
22629	52997	A	22757	<u> </u> 1	2127	MDQSGLLALRAAVGAVVGSTL
22029	3.2337	-				AVDIGILVKDWHPRPLGGKHC
ļ						QLLGAFRCRQHRRGKEERRGE
						TKEKKGERENERGKRRRIEGER
		İ				DKYVKENEKQTCVQKERHSCT
						GNSPNHMATVQAERGALDSHG
l						GWHPWDYCAVQQGRYHQLLL
						TKEEIEVQRSKKMCLLQAFLIC
						CQKSHFLGPGCTRSMVPASASG
						ESLKLLPLMVEDEGEPVTERE\R
						KREGERERGRERERRRRKKK
						RRRKRRKRRRRRRRRG*EFLS
						LG*NERGKRRRIEGERDKYVKE
						NEKQTCVQKERHSCTGNSPNH
		1				MATVQAERGALDSHGGWHPW
1						DYCAVQQGRYHQLLLTKEEIE
						VQRSKKMCLLQAFLICCQKSHF
						LGPGCTRSMVPASASGESLKLL
		ŀ				PLMVEDEGEPVTEREQEEGGGE
		l				RERERERKKEKEKEKEKEKEEK
		l			İ	EKEKEKEEERIRVPLPRVELRLT
		l				NIAAKQVGNTILDVAAPADHPS
						ESSCVYVRFVIKIPVAASFRFKK
Ì						SYVTNNVESCEDGIHYFWKSH
1						MHSENSCSCKLPCPPHLPAAER
			Ì			LGDALEPLLPTMRRQRQCQNT
			ł			GDCDGRRQIGTRETEMHRDTSL
		1				VAVKTQVDTSREGMGEVEPNS
						CFWRDTHMASGVGAAFEELPH
						DGTCDECEPDEAPGAEEVCREC
						GFCYCRRHAEAHRQKFLSHHL
						AEYVHGSQAWTPPADGEGAGK
22630	52998	A	22758	348	881	WRQPRWPHHQACLWQRAGSS
		Ì				YSQHKDPHFTSFSTPRAVQNFD
						GFPCPFLRETTTLVLIPSVPLTDP
						GSFYRAWSFQCGPRSPRPLLGR
		ļ				TLSQKFLLRLRGRPRRPYMSTL
		1				TPF*STRIFERAFCQVMGAESGG
						RIEKTTLCKRSLGPLMEPRV
22631	52999	Α	22759	1	347	MVPASASGESLKLLPLMVEDER
						EPVTERE/REKGRGREREGERE
					1	KEREGERERERGERKGERER
				1		RRG*QPPFHNRFSEGQGRSFLCS
						RRPRVSRRGHTSRGPARVKAAE
		}			<u> </u>	RRGLG
22632	53000	Α	22760	145	258	PENGLIGLS*KIVLSLNIPVNEN
		•				DMKVDEVLYEDSSTA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22633	53001	A	22761	1	1479	MALKEIFCPVRETETCTINCGTR
						ISPSILHVGLPYPSTPNVKPERE
						GCGGGVSGNSKQPGALETAESI
						GKKEGRREEGSRPESASSLSLLL
						SLAYDVDKSAQICTRIHSSVPLE
						AVYPGYIHQSIINQADLSALAPY
						DPCVCQIHKYHGAYNCLQYSV
						QNTLYSPTTEQYKRCQAPWVL
		1				PAIFTIPASGAAASAVPVRAQA
						QGGQAAGQRRRLATAPLTGEG
			}			EPDHGDRTRAGSQGPEARPRIH
						LGAARGGDRTRGRPGRRGRDW
						TRLGREMAAAAVSSAKRSLRG
						ELKQRLRAMSAEERLRQSRVLS
						QK\VIAHSEYQKSKRISIFLSMQ
ļ		Ì				DEIETEEIIKD\ICQRGK\ICFIPPV
						RFSRAITMDMVRIGSP\EGNFFN
ŀ						FPKTSWEYSLRLGEG\DVREEG
		ŀ				LVPPGGL/EILILPCPQALGLDQ
						NHGQPRLG\RGQGAYL*CLS*K
[RLFCSHSGKLKPPTPPGGWLFK
		1				EQICPPRSPVNENDMK\VDEVL
22634	53002	Α	22762	1	813	
22635	53003	Α	22763	1	368	
22636	53004	A	22764	327	4281	SASASRCDPGSGSRREREEELQ
						WRRRRRRRRRRRRRQRQRA
						AAPAAPAGGIEAVNMASASYHI
						SNLLEKMTSSDKDFRFMATND
						LMTELQKDSIKLDDDSERKVV
						KMILKLLEDKNGEVQNLAVKC
						LGPLVSKVKEYQVETIVDTLCT
						NMLSDKEQLRDISSIGLKTVIGE
						LPPASSGSALAANVCKKITGRL
	İ					TSAIAKQEDVSVQLEALDIMAD
						MLSRQGGLLVNFHPSILTCLLP
						QLTSPRLAVRKRTIIA
22637	53005	A	22765	2	271	
22638	53006	Α	22766	299	451	IVALFRSLVRHLRGTMAMESTA
						TAAVAAEL/VSADKIEDAPAPST
						SADKVES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amîno acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22639	53007	Α	22767	1	1122	MATESTATAAIAAELVSANKIE
22033	33007	ľ`	22707		1122	DVPAPSTSADKVERIENGVLGN
						TLDGVHVEEEEGEKTEDESLVE
						NNDSIDEEARKELREQVYDSM
						GEKEEAKKTEDKSFAKPETDKE
						QDCEMEKGGREDMDISESAEEP
						QEKVDLTLGWLTEISEGAKGG
						GAPEGPNEAEVTSGKPEQEVPD
		İ				VEEEKSVSETDVQKECREKGG
						WEKHREVIVSIEEKPKEVSEEQP
						VVILEKQGTAVEVAAESLDPTV
ļ						KPVDVGGDEPEEK/GTVPGSLR
						PSPCRDPLPAGLGLWVQLSV**
		1				GSGTVQQIY*SH*EENGCTKRA
		l				GEGG*RIVC*IQERN*GTEGTAT
						RNYRKDRRCKGVSDYWECS*T
		l				GSESYSGGELYFRFHS*WRRLF
22640	53008	Α	22768	223	339	OSESTI OGESTI IGI IIS WIGGI
22641	53009	В	22769	341	496	
22642	53010	Ā	22770	12	1099	VSGVPNCLFFSQALFRSLVRHL
22012	33010	`		1.2		RGTMAMESTATAAVAAELVSA
						DKIEDVPAP\STS\ADKVESLDV
					Ì	DSEAKKLLGLGQK\HLVMGDIP
						AAVNAFQEAASLLGKKYGETA
		ŀ				NECGEAFFFYG\KSLLELARME
						NGVLGNA\LEGVHVEEEEGEKT
						EDESLVENNDNI/D*N*RLGRG*
		ļ				*RK**D*RNAK*FSP*KQVSSRK
	1					*GGGDWEPRACLGYAGFSKDH
						F*KARNKRSTALCCPGTS*TRRS
				i		*C*I*\KLCASC\GGVPVLP*PTGP
						VPGSPRPSPCRDPLPAGLGLWV
						QLSV**GSGTVQQIY*SH*EQNG
						CTKRAGEGG*RIVC*IQERN*GT
						KGTATRN*REDRRCKGVSA*W
						ECS*TGSESYS
		1	l			ECO TOOESTO

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22643	53011	Α	22771	11	2485	VSGVPNCLFFSQALFRSLVRHL
						RGTMAMESTATAAVAADVVS
						ADKIEDVPAPSTSADKVESLDV
						DSEAKKLLGLGQKHLVMGGIP
						AAV\NAFQGAA\SLL\GKKYGET
						ANE\CGEAFFFYG\KSLLELARM
						ENGVLGNALEGVHVEEEEG\EK
	ĺ		:			TEDESLVENNDN\IDEEAREELR
						EQVYDAMGEKEEAKKTEDKSL
						AKPETDKEQDSEMEKGGREDM
	-					DISKSAEEPQEKVDLTLDWLTE
						TSEEAKGGAAPEGPNEAEVTSG
					·	KPGQ\EVPDAEEEKSVSGTDVQ
						EECREKGGQEKQGEVIVSIEEK
						PKEVSEEQPVVTLEKQGTAVEV
						EAESLDPTVKPVDVGGDEPEEK
						VVTSENEAGKAVLEQLVGQEV
						PPAEESPEVQTEAAEASAVEAG
			,			SEVSEKPGQEAPVLPKDGAVN
						GPSVVGDQTPIEPQTSIERLTET
						KDGSGLEEKVRAKLVPSQEETK
						LSVEESEAAGDGVDTKVAQGA
						TEKSPEDKVQIAANEETQEREE
						QMKEGEETEGSEEDDKENDKT
						EEMPNDSVLENKSLQENEEEEI
						GNLELAWDMLDLAKIIFKRQET
						KEAQLYAAQAHLKLGEVSVES
						ENYVQAVEEFQSCLNLQEQYL
						EAHDRLLAETHYQLGLAYGYN
						SQYDEAVAQFSKSIEVIENRMA
						VLNEQVKEAEGSSAEYKKEIEE
						LKELLP\EIREKIENAKE\SQR\SG
22644	53012	Α	22772	52	161	FQNPTGLY/CDTEYN\RLQKQQ
22044	33012	1	22112	32	101	TEKTKQKKTQFHPICN
22645	53013	Α	22773	3	115	FFQNPTGLY/CDTEYN\RLQKQQ
22043	33013	7.	227,3]		TEKTKQKKTQFHPICN
22646	53014	A	22774	229	358	TEATRONKI QI III ION
22647	53014	A	22775	257	386	
22648	53015	A	22776	259	294	
22649	53017	A	22777	217	398	KAWWNLLRKSEPKSCGS*GGL
22049	2301/	^ <u>`</u>	22///	21/	370	STSRERLVECSSPLGLGYCG**L
						CHLQEPHYGSLHRMSS
22650	53018	A	22778	1	3386	CHECETHOSPHONOS
22030	155018	I^1	22//0	<u> </u>	3300	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
}		1		sequence		
22651	53019	A	22779	391	2100	SRRTAGVRLDILCKDPIMWVSA
22001						DGHSYQPQYPKAKRATEFHFF
						NFKALLARAAVGAARGIHIHRC
				ť		GLTKRPDARRRRVAGAQGREP
1						AISLPGDRAAGARATRTRGPGP
		ĺ				APKMPAIAVLAAAAAA\WCFL
1						QVESRHLDALAGGAGPNHG\NF
ļ						L\DNDQ\WLSTVSQYDRDKY\W
 :						NRFRDEVEDDYFRNW\NPNKPF
]			DQAL\DPSKDPCLKVKCSPHKV\
						CVTQ\DYQTAL\CV\$RKHLLPR
1						QKEWGTWAQKHWGLDL*ILV
						K\CKPCSRGRSSAMGLAGSDGP
						LLTAFQVANLEFHACSTGQK/S
						FATLCDG\PCPCSSQSLEPPKHK
						GRKGVPCTDKELRNLASRLKD
						WFGALHEDANRVIKPTSSNTAQ
						GRFDTSILPICKDSLGWMFNKL
		ļ				DMNYDLLLDPSEINAIYLDKYE
						PCIKPLFNSCDSFKDGKPFLNNE
						WCLLPSQNPGGLP/CAQNEMNR
		1				IQ\KLSKGKSLLGAFIPRCNEEG
						YYKATQCHGSTGQCWCVDKY
						GNELAGSRKQGAVSCEEEQETS
						GDFGSGGSVVLLDDLEYERELG
						PKDKEGKLRVHTRAVTEDDED
						EDDDKEDEVGYIW
22652	53020	Α	22780	2	441	HGVQGAGCGEVAAVSIELGAG
		l				GPVWRAAPRACSRLQLLRGFG
		l				LEGQWFGESCCRRGAS*EGSTR
						*PPPRRQPWPEWVMSWVPPET
-						REPARISPAPAKQN*PPPAPKSP
						AAPA\PQAAVAATPSPPGEISEA
						WDSLESFSKPAFFSQ
22653	53021	Α	22781	3	121	

SEQ ID	_		SEQ ID NO:			Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>				
22654	53022	A	22782	2	1488	FVNNMADGEEPEKKRRIEELL
						AEKMAVDGGCGDTETWKDR\
	,					WNHVKK\FLERSGPFTHPDFEP\
						STESLQFLLDTCK\VLVIGAGG\
						LGCELLKNLALS\GFRQIHVID\
						MDTI\DVSNS*IRQFLFRPKDIGR
						PKAEVAAEFLNDRV\PNC\NVV
		1				PHF\NK\IQDFN\DTFY\RQFHI\IV
						CGTGTLSCARRWINGMLISL\LN
						YERMVS*DPSSIVPLID\GGTEGF
						\KENARVIL\PGMTACIECTLEL
						YPPQVNFPMCTIASMPRLPEHCI
						\EYVGMLQWPKEQPFGEG\VPL
						DGDDP\EHIQWIFQKSLERASQY
						NIRGVTYRLTQGVVKRIIPAVA
						STNAVIAAVCAT*/EVFKIATSA
						YIPL*IITWVFNDV\DGLYTYTFE
						GRKGK\ENCPAC\SQLPQNISVF
						LHQAKLQ\EVLGIILTNSASLQN
		İ				EILPAITATLGGEKIGTLYLQSV
						TSI\EERNRANFFQTIERNWGLL
		1				DGPKTGRLLDVTTP\QTVLFQTS
						ILLLKGKSPT
22655	53023	Α	22783	1	1719	
22656	53024	В	22784	199	460	
22657	53025	Α	22785	226	1620	NGSGLALATGNRDPGLCPLSAP
İ						SCPWR*KSAGASTQQAYSSQSQ
		İ				/HPAYHRITPT\LTNFALRLYKEL
						AADAPGNIFFSPVSIS\TTLCLLC
						LG\AQANNSALILEGLGFNLTET
		ŀ				PEADIHQGFRSLLHTLVLPSPKL
						ELKVGNSLFLDKRLKPRQHYFD
						SIKELYRAFAFSAHFTDSVTTGR
						QINDYL*RRTYWPGVDLPFRS/L
		l				ARDTFMVLA\NFILFSKAKVKA
		1				TLSSRVPDPKSRPVFLVDERTSF
						\QVPMMHQKEMHRFLYDQDLA
						CTVLQIE\YRGNALALLVLPDPG
						KMKQVEAALQPQTLRKWGQL
						LLPSLLDLHLPRFSISGTYNLEDI
						LSPNWFSPTYTLNKITSLQSVG
						G*LNKSISKVSHKSMVDMNEK
						GDQRPGLLQASSPSPHL*/YTMS
						DPHAHFNRPFLLLLWEVTTQSL
						LFLGKVVNPVVWVTMVGGQE
						LSYLILDQTDRPEPACILGLLCG
22658	53026	A_	22786	64	288	
22659	53027	Α	22787	1	1923	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22660	53028	Α	22788	309	2286	CAVCHISFQDTPVLVSSNVTMQ
						FGSKPLFENISVKFGGGNRYGL
						IGANGSGKSTFMKILGGDLEPT
						LGNGSLDPNER\IGKLRQ\DQFA
						FEEFTVLDTVIMGHKEL\WKVK
		ĺ				QERDRIYALPEMSEEDGYKVA
						DLEVKYGEMAGYSAEARAGEL
						LLG\VGIPVEQHYGPMSEVAPG
ĺ						WKLRVLLAQALFADPDILLLDE
						PTNNLDIDTIRWLEQVLNENNS
						T\MIIISHDRHFLNMVCTHMAD
				1		LDYGELRVYPGNYDEYMTAAT
						QARERLLADNAKKKAQIAELQ
				1		SFIDKIKLEEVKASSRQNPFIRFE
						QDKKLFRNALEVEGLTKGFDN
						GPLFKNLNLLLEVGEKLAVLGT
						NGVGKSTL\LKTLVGNLQPDSG
						TVKWSENARIGYYAQDHEYEF
·						ENDLPVFEWMSQCKQ*GDDEQ
						AVRSIFGRLLFSQDVIKKPAKVL
						SAPFCKDSILEAIDAGIKLIITITE
	1					GIPTLDMLTVKVKLDEAGVRMI
						GPNCPGVITPGECKIGIQPGHIH
						KPGTGELNITTGGIVKARDTQIA
		ŀ				LNDKSKGDVRVDGQNSLLETF
		l				NMYVGTSGTGTLTLTNNGTLN
						VEGGEVYLGVFEPAVGTLNIGA
						AHGEAAADAGFITNATKVEFG
						LGEGVFVFNHTNNSDAGYQVD
22661	52020	ļ.	22790	1	552	MLITGDDKDGKVIHDARHTVF
22661 22662	53029	A	22789 22790	1633	553 2207	ILFFDKGMPAQTPSLQKRWEK
22002	33030	A .	22/90	1033	2207	M*LVLW*GDREGIKAI/ISEESW
						GARAPIFLLR/MMEDQGLYSIR
						MGGARGLPGKSLAL*GAGRGF
			•			YSGERTWTLSLPSGNCAP\YQR
						GGWWYHACA\HSNLNGVWHH
		1				GGHYRSRYQDGVYWAEFRGG AYSLRKAAMPHLAPEAVTLCSF
		1				GPLGPRGHWSAGAPSCSGHTFF
22663	52021	A	22701	34	210	AAADPLRHLRRGAFIKAGRHA
22003	53031	^	22791	34	210	
		1				G/PSLPVCLQATA\H*MDLARLA
L			L	<u> </u>		LVLVVRTSASTKSSTRE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22664	53032	Α	22792	397	1168	LLSRISKVSSKLPVRGGAGSLCT
						GDQRK\RRKFL\ETVELQISLKN
						YDPQEGQSAFSGHPSRL*V\PSP
	1					PLSSLVCVLGGPSKHL*RRLRP
	-					VGYPPTMDIEGA*KKLNQELKN
]				LVKKAGPRKYDCVFWPQESLD
		1				SSKIPPNPSGPGFK*RAGKVPFP
		1				CLTHNGKHGLAQSGIEVEVPTF/
						LSFQMKKGVYVWAVAVWSRE
						R*QTNELVYNIHLACQLSWVSL
						LKEKNWQ\NVPGPYIIKSTMGQ
		_				SPSSHILRHIRINSITS
22665	53033	Α	22793	1	377	SVSRVPTRSLRTSTWRLAQDQT
		1				QDTQLITVDEKLDITTLTGVPEE
						HIKTRKVRIFVPARNNMQSGVN
						NTKKWKMEFDTRERWENPLM
						GWAST\WSYDIEERKVPKPKSK
						SYGANFSWNKRTRVSTK
22666	53034	Α	22794	183	368	
22667	53035	Α	22795	3	623	ILAFACSKMAAVSMSV\VLRQT
						LWRRRAVAVAAL\SVSRVPT/R
						RSLRTSTWR\LAH\DQTQDTQLI
						TS**KIGISLTFNWPFPEEHIKNL
			1			EKSGIFVPCSPITCQSGVNNTK\
			1			KWK\MEFDT\RER\WEN\PLMG\
						WASTGLIPLS\NMVLNLPVLKE
						DAVS\FAEK\NGWELWTLEERK\
						VPKPKSQVFMGANFS\WNKRTR
						VSTKIGLALTISPA
22668	53036	Α	22796	1	798	MNKVNPTQQDKEDHPELIITSIG
	1					DKRQLTLEAASGVCHRGTQTR
						VKEESKKFLISMSFRGGKPVHK
					†	VVERACVCTCLHHTSVPATHN
						TQRRHTTPTTRKNGRKRRGKK
						EREKRKDREHNTGRRTKRKKT
						RNIGGQRTQQKQKQSEDKARR
		1				EPGGGRKQGRGREDGTRQRRG
			-			GRTRNREEGAQRGRNGR/RGEG
						RRTEEETHMA*APDITATQQTE
						QQGHGTRQRDTTLRRHTAEDA
						QPSEHASADQAHTKKTE*SVSR
		L.				TNGQYLLT
22669	53037	В	22797	85	322	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22670	53038	A	22798	165	4733	AATALLTHGQEEGQVEGQDED IPPITCVQNGLRYHDRDVWKPE PCRICVCDNGKVLCDDVICDET KNCPGAEVPEGECCPVCPDGSR VTHRPKKPPGVEGPKGDTGPPR PKGT\PGPPGRDGIPGQPGLPGP PGPPGPPGPPGLGGKLLLPKLSY GYDEKSTGGISVPGPMGPSGPR GLPGPPGAPVSIQDVFICLLGLW SFGGKTGMRAGEMLRDLLGPQ GFQGPPGEPGEPGASGPMGPRG
						PPGPPGKNGDDG
22671	53039	С	22799	88	392	
22672	53040	Α	22800	37	293	
22673	53041	A	22801	55	1917	GGGGVRRRRGTRRSEPGDSRSA MAPIGLKAVVGEKIMHDVIKK VKKKGEWKVLVVDQLSMRML SSCCKMTDIMTEGITIVEDINKR REPLPSLEAVYLITPSEKSVHSLI SDFKDPPTAKYRAAHVFFTDSC PDALFNELVKSRAAKVIKTLTEI NIAFLPYESQVYSLDSADSFQSF YSPHKAQMKNPILERLAEQIAT LCAT\LKEYPA\VRYRGEYKDN ALLAQLIQDKLDAYKADDPTM GEGPDKARSQLLILDRGFDPSSP VLP*I*LFQGMSY*SCLPI\ENDV\ YKYETSGIGGGHGVKGGASGT EDDDLW\IALR\HKHIAEV\SQ\E VTRSLKDFSSSKRMNTGEKTT MRDLSQMLKKMPQYQKELSK YSTHLHLAEDCMKHYQGTVDK LCRVEQDLAMGTDAEGEKIKD PMRAIVPILLDANVSTYDKIRIIL LYIFLKNGITEENLNKLIQHAQI PPEDSEIITNMAHLGVPIVTDST LRRRSKPERKERISEQTYQLSR WTPIIKDIMEDTIEDKLDNK\HY P\YISTRSSASFSTTAV\SARYGH WP*GTRPPGEYRSGP/RVFIIFHP LGGVSLNEMRCALRR*PRANG KWGGA**GSTHILQPHRNLPGH

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SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22674	53042	A	22802	46	1931	GGGGVRMLCGTRRSEPGDSRS AMAPIGLIAVVGEKIMHDVMI. NDLNMGEWKALLVDQLSMRM LSYCCLMTDIMTEGITIVEDINK RR*PLPSLEAVYLITPSEKSVHS LISDFKDPPTAKYRAAHVFFTD SCPDALFNELVKSRAAKVIKTL TEINIAFLPYESRVYSLDSADSF QSFYSPHKAQMKNPILERLAEQ IATLCATLKEYPAVRYRGEYKD NALLAQLIQDKLDAYKADDPT MGEGPDKARSQLLILDRGFDPS SPVLHELTFQAMSYDLLPIEND VYKYETSGIGEARVKEVLLDED DDLWIALRHKHIAEVSQEVTRS LKDFSSSKRMNTGEKTTMRDL SQMLKKMPQYQKELSKYSTHL APCLRTCMKHYQGTVDKL\CR V\EQDLAMG\TDAEG\EKI\KDP MRAIVPILLDANVQTY*QIRIILL YIFLKNGITEENLNKLIQHAQIP PEDSEIITNMAHLGVPIVTDSTL RRRSKPERKERISEQTYQLSRW TPIIKDIMEDTIEDKLDTKHYPYI STRSSASFSTTAVSARYGHWHK
						NKAPGEYRSGPRLIIFILGGVSL NEMRCAYEVTQANGKWEVLIG STHILTPTKFLMDLRHPDFRESS RVSFEDQAPTME
22675 22676	53043 53044	A	22803 22804	2 236	51 451	LHLPLFLGAIRSGQYGLWNLFR VLGWTCSLGIYLGWRVS*NFG RGTKALKVLSTIS/E*RNHHCHP *NNIFVY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22677	53045	Α	22805	3	1173	GAGAPAAGQAGLTSTSAVRAR
						PVSPARQSQDPAASLSLHLSLPP
						APGAGCARWRRRGPQAAMAA
						GAGAGSAPRWLRALSEPLSAA
						OLRRLEEHRYSAAGVSLLEPPL
						QLYWTWLLQWIPLWMAPNSIT
		l				LLGLAVNVVTTLVLISYCLTAT
		İ				EEAPYWTYLLCALGLFIYQPLD
						AIDGKQARRTNSCSPLGELFDH
		1				GCDSLSTVFMAVGASIAARLGT
	j	l				YPDWFFSRSFYLGCLVFYC\AH
		1				WQTYVS\GMLRFGKVDVTEIQI
						AL\VIVFVL\SAFG\GATM/WGD
ļ						YTIPILEIKLKILPVLGFLGGVIFS
						CS\NYSHVILH\GGVGKNG\STIA
						GTSVLSPGTPH/MGLIIIL\AIMIY
	ŀ	İ				*KVRQLMVF*KSIPCL*YPQCLG
						CVF\AKVSQKISGSSHDQK
22678	53046	Α	22806	2	1091	CPGRRRRRRPGRSRARATALPT
	ļ					PGPRKRGGPAGRKRERPERCSS
		1		•		SSGGSSSSGDEDGLELDGAPGR
				:		GKRAAQPVAAGKVGGAAVVV
]				AEPEHTKKHVKLETLLITTEGK
						LWSWGRNEKAQLGHGDTKRV
	}					EAPRLTEGLSHEVIVC\TCARNH
	ŀ					TLALTETGSVFAFGENKMGQL
						GLGN*TDAVPSPVQIMYNSLPIT
						KMA/CGYGWLGYTEQKDEMV
1						PRLVKLFDFPGHGASQIYAGYT
						CSFAVSEVGGLFFWGATNTSHE
						STMYPKAVQDLCSWRIWSLAC
					1	GKSSIIVAADESTISWGPSPTFG
						ELGYRDHKPKSSTAAQEVKTL
						HGIFSEPVAMGYSHSLVIARDE
						SETEKEKIKKLPEYSPQTL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22679	53047	A	22807		1409	MSLGLASERLWAGLGLPALPL ADEGGECGETLFVWRSQKLEG SKCKGQLLIFGATNWDLIGRKE VPKQQAAYRNLGQNLWGPHR YGCLAGVRVRTVVSGSCAAHS LLITTEGKLWSWGRNEKGQLG HGDTKRVEAPRLIEGLSHEVIVS AACGRNHTLALTETGSVFAFGE NKMGQLGLGNQTDAVPSPAQI MYNGQPITKMACGAEFSMIMG LQRKPLFLWGALNMGQLGTQL RMGKFHSPGAQRIEYDCELVPR RVAIFIEKTKDGQILPVPNVVVR DVACGANHTLVLDSQKRVFSW GFGGYGRLGHAEQKDEMVPRL VKLFDFPGRGASQI/SCWL/HTC SLCCQ*NGWSVFLGGHQHLP*I YHVPQKQCRTSAAGESGTLAC GKSSIIVAADESTISWGPSPTFG ELGYGDHKPKSSTAAQEVKTL DGIFSEQVAMGYSHSLVIARDE SETEKEKIKKLPEYNPRTL
22680	53048	С	22808	277	518	
22681	53049	Α	22809	2	192	
22682	53050	A	22810	258	2946	MAEAGLRGWLLWALLERLAQ SEPYTTIHQPGYCAFYDECGKN PELSGSLMTLSNVSCLSNTPAR KITGDHLILLQKICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAGQLPAV VAYEAFYQHSFAEQSYDSCSRV RVPAAATLAVGTMCGVYGSAL CNAQRWLNFQGDTGNGLAPLD ITFHLLEPGQAVGSGIQPLNEGV ARCNESQGDDVATCS
22683	53051	A_	22811	258	362	
22684	53052	A	22812		4038	MAEAGLRGWLLWALLLRLAQ SEPYTTIHQPGYCAFYDECGKN PELSGSLMTLSNVSCLSNTPAR KITGDHLILLQKICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAGQLPAV VAYEAFYQHSFAEQSYDSCSRV RVPAAATLAVGTMCGVYGSAL CNAQRWLNFQGDTGNGLAPLD ITFHLLEPGQAVGSGIQPLNEGV ARCNESQGDDVATCS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22685	53053	Α	22813	198	380	ASSESDSTPCR*MAYSWPPNSN
						SSHCIKELPLFMGKIPAAS*DGG
						PQGLAIFPPGVIVGI
22686	53054	Α	22814	1	154	
22687	53055	A	22815	35	486	PARRGRDEC*VLAPFRATRPPS
						RRTR*SKATYPTPSNDQTRTIRT
						SRERASMPRSAEPKRPTIPRLHP
						SPRGIVTVSS/RPRSHALPVSHIH
						SCHSRTAISDNQVPAFLQVYGE
		ŀ				EVRKRAGEVTQAITGFPRYRLF
						FKMLRCHRSIQVTAST
22688	53056	Α	22816	212	365	RCRSGPAPRSAPGARPASPSPPA
						AGTCR*RPAGHCPAGAAGLLSL
						CPRCST
22689	53057	Α	22817	293	618	QQNPTNCLPPASR*SKT*RARST
						AGPTRPHLHHPAAPAPGSPACA
						GEGRLQR/HSPNPLYFNWKKQR
						FAAKPHFRRPQSSLNPQQWLLA
						GLLQEESSSLGTDTAPLHSLG
22690	53058	Α	22818	3	1129	FCLDDLSSAINEHLSARGRRRRS
						IVKVSLHPAVIGLKSKFLKKPD
						QLRKLF\IGGLSFETTDESLRSHF
						EQ\WGTLTDCVVMRDPNTKRS
						RG\FGFVTYATVEEVDAAMNA
						RPHKVD\GRVV\EPKRAVSREDS
						QRPGA\HLTV\KKIFVGGIKEDT
						EEHHLRDYFEEIILNSMGKLEV
						VEIH*LDR\GQWPRKRGFAFVT
						FDGHDSVDKIVIQKYHTVNG\H
						NCEVRKALSKQEMASASS\SQR
					}	G\RVGSGNFWCGRT\NGFGGND
						NFG\RGGNFSWSVVAFGG\SRG
						GVWIWVAVGDGYNGFG\NDGS
						HFG\GGGSY\NDFGDFH\NQSSK
						FWTP*RGGNLGGQKLWPPMGG
						GG\QYFAKPTKPKVAYGRFPAA
						ASSYGK WPEDFN
22691	53059	A	22819	1	568	MGKMSPGHVRDFHGSCHHNRP
						GGPGGKSGFMGQALGPG/GCV
						QPRDLVLCVSATPAMAERGQC
						GAQAVASEGGSPRPWQLPCGV
						EPADFRRYMETPGCPGRNL/PA
						GAGPL*RTSAGAVWKGNVRSE
						SP/PQSLYWGTASGAVRRGPPSS
						RNQNGRSTDSLHCMPGKDTQ\P
						NP*KQPGGRLHPAKPQGRSCP
22692	53060	A	22820	232	389	, , , , , , , , , , , , , , , , , , , ,
22072	122000	111		232	1557	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
i				sequence		
22693	53061	Α	22821	3	401	VGVCEKRGAPG\PPLAPGPARG
						PVAAAVPP/GPGASRARP*DAA
						AAAGGRGPPEQ*T*KAWGLHP
						GTGSL/RGHPAPM*PLLRAFPRS
						PLHPLPPPHPLPRRAA\SNPGTS
-						KCRTAAGTRQWGGRQSQLQRC
						QCLR
22694	53062	Α	22822	1	410	MGKMSPGHVRDFHGRPSHHRP
						RGPGGKSGFVGKVQGPHAVCS
						LGTWCLVSQPLHP\G*KGPMYS
		l				SG\VASEGGSPKPWQLPRGVEP
						MGAQKSRTEVWEPLPRFQKIY
						GNTWMPRIPMCCARDPGGAKG
						GCSNPSQRR
22695	53063	Α	22823	264	979	GPRHADPANFSARNLSRP*DAP
						GRVQDPRPQAPQL\PPGLGAPSR
						VRRPPPRRPVDRTLPGGPRPPA
						AAA\GPTGGRGS\PRAGGTAAA
						TGPRAGPGARGAAGSAPFLTDP
						NM\GRPPPPATTTPPGAATAPGP
						REEAPGKFRVSPPGPSGAGGPP
		1				TRASPTPTAAGGGGGDSEGEGP
1						ILLVADAG\PDSARSPPGLPGCV
ł		ŀ		<u> </u>		LSAAPRSRRDPDAVGGLAASA
			<u></u>			AARRLNTTWKLPQLGASTL
22696	53064	A	22824	271	426	
22697	53065	Α	22825	546	1296	KMVDCLANSEANTRRISIVENC
1		l				FGAAGQPLTIPGRVLIGEGVLT
						KL\CRKKPKARQFFLRNDIVV*
		İ				GNIVIQKKKYIKQHIIPLENVTID
						SIKDEGDLRNGWVIKTPTKSF\A
		Į.				VYAATATEKSEWMNHINKCVT
İ						DLLSKSGKTPSNEHAAVWVPD
		1				SEATVCMRCQKAKFTPVNRRH
		l				HCRKCGFVVCGPCSEKRFLLPS QSSKPVRICDFCYDLLSAGDMA
		1				1 ~
						TCQPARSDSYSQSLKSPLNDMS DDDDDDDDMQ
22698	53066	A	22826	1219	2370	МОПОПОПОМО
22699	53067	A	22827	164	601	DRTNIMVGYHQTNQKTDTGKT
22033	33007		22021	1107		PTRRPVLVDH\NRLPEGSRGRL
						AVAVAGDHPAAVQVTMTLVN
						DTGFDPVFSGSIAESWGQHSGT
						PSYC*DWEAATMLRAFPLAKK
						GETKDLRQFPNIDNAYMELGT
						NRADAVLHDTPNILYFMQF
22700	53068	A	22828	2	91	THE PARTY DELICATION OF THE PARTY OF THE PAR
22,00	10000	l' • -	122020	<u> </u>	I *	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22701	53069	A	22829		1188	MGAEPSLLTSLGTLWSGERFQC MRAKIWERSRKGRRFSTDVISP TKDSFAGPFQPISDTLVQHQRIY GLKVCVLPDPCKIRILKPNPQC VGKWEEEVAENEIQVFGGERF DINKYSKSPPPPTPQPLYLTAGF ASCRNHRKSEELREGKGESPGK SKPFHLQANFTVALVTTLQGDK HSQEAFIRPVAKAKTTCTYLSN VCEMKDFKRINCPDPFHKMAP KAKKEA\PAPP\KAEAKAKGFK GPRRQVLK\GVHNHKKRKKK ERSRTFTHLSRRPGRPLRI\RRQP \KYPSGRAAPRRNKLD\HYAIIK FPLT\TESA\MKKIEDNNTL\VFI\ VELLKATTKNQIKQGC*RSLYD I\DVGQGSTPWISAWMGEKKGI CSQLAPDLPIAFGMFANKIWGFI
22702	53070	Α	22830	1413	2197	RVEWRWYNVLPCLDVHSGPVD RRSHVMTAAEKPPFCPAWGQA VSQQSCALEPRTGEEGGKGIQG PASGLFPTGEEISVVPNRWLDA SLSPVILGVQG\GSQCLSCGVGQ EPTLTLEVRLGASSLGTQPQML SLLKPGSPQPWCCGTP*QDSVD GSFASSLRILAKALLCGPGFCPS LTLPSFCRKPVTIME\LYLGAKE S\KSFTFYRRDMG\LT\SSFESAA\ YPGWFLCTEPE\ADQPVRLTQL PENGGWNAPITDFYFQQ\CD
22703	53071	A	22831	100	177	KRDHF*SRLQQVSRSVRWSSSA TDSR
22704	53072	A	22832	610	1080	
22705	53073	Α	22833	1169	1308	

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SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22706	53074	A	22834		1889	ACRRAGTWTPPRPRGSGSRA MAGNCGARGALSAHTLLFDLP PALLGELCAVLDSCDGALGWR GLAERLSSSWLDVRHIEKYVDQ GKSGTRELLWSWAQKNKTIGD LLQVLQEMGHRRAIHLITNYGA VLSPSEKSYQEGGFPNILFKETA NVTVDNVLIPEHNEKGVLLKSS ISFQNIIE\GTRNFHKDFLIGERG E/IFGGYQEGKIQTLTYAVKLFK QEKKMQCKKHWKRFLSELEVL LLFHHPNILELAAYFTET\EKFC LIYPYMRNGTLFDRLQCVGDT APLPWHIRIGILIGISKTIHYLHN VQPCSVICGSISSANILL\DDQFH PKLTDFSMAHFRSP\LEHQSCCI NMPSS\SRKHLWY\MPEEYIRQ GKLSIKTDVYSFGIVIMEVLTGC RVVLDDPKHIQLRDLLRELMEK RGLDSCLSFLDKKVPPCPRNFS AKLFCLAGRCAATRAKLRPSM DE\V*RALESTQASLYF\AGDPP TSLKALRGP\SPLF\LEKVT\SIPV \EDDE\KPNNNLLPS**KALRMK RWPQK\TPF\ECSQP\EVWFLSLE QKAREARENEEACNMPSSS\CE
						ESWFPKY\IVPS\QDLRPL*GK*K ILSSQA\SGHSC\RTKPVESSCSS KFSWDEYEQYKKE

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22707	53075	A	22835	1	1560	RWERGGKRFRAADPMSALRRS
				-		GYGPSDGPSYGRYYGPGGGDV
		ŀ				PVHPPPPLYPLRPEPPQPPISWR
						VRGGPAETTWLGEGGGGDG
						YYPSGGAWPEPGRAGGSHQEQ
		ĺ				PPYPSYNSNYWNSTARSRAPYP
						STYPVRPE\LQGQSLNSYTNGA
						YGPTYPPGPGANTASYSGAYY
						APGYTQTSYSTEVPSTYRSSGN
		1				SPTPVSRGIYPQQDCQTEAPPLR
						GQVPGYPPSQNPGMTLPHYPY
						GDGNRSVPQSGPTVRPQEDAW
		l				ASPGAYGMGGRYPWPSSAPSA
			1			PPGNLYMTESTSPWPSSGSPQSP
						PSPPVQQPKDSSYPYSQSDQSM
		İ				NRHNFP/CASVHQYESLGDQ*T
						MDDSRILLDSPSPSIVPEPSACM
			1			GNAHPVTHP\NNQDQSSSLPEE
						CVPSDESTPPSIKKIIHVLEKVQ
						YLEQEVEEFVGKKTDKAYWLL
						EEMLTKELLELDSVETGGQDSV
		l				RQARKEAVCKIQAILEKLEKKR
						IMKGFITKWKPVTNLTKEHLIW
						LITLFLKCLLMTRSNTFQLFL
22708	53076	A	22836	8	207	TPVPRALRSPAGLEGSHLSSIST
22700	33070	``	22030	ľ	207	MMPAVMSVGKVTENGGSPQGI
						KSPSKPPGPNRIG*RNQVSVHT
22709	53077	A	22837	459	5520	RQVRPAGLEGSDLSSINTMMSA
22,05	33077	 	22037		3320	VMSVGKVTENGGSPQGIKSPSK
		ŀ				PPGPNRIGRRNQETKEEKSSYN
						CPLCEKICTTQHQLTMHIRQHN
		l				TDTGGADHSCSICGKSLSSASSL
						DRHMLVHSGERPYKCTVCGQS
						FTTNGNMHRHMKIHEKDPTSA
;						TATAPGRGDFASAKSSKRKLSH
		l				DAESEREDPAPAKKMVEDGQS
		ŀ				GDLEKKADEVFHCPVCFKEFV
		l	ļ			CKYGLETHMETHSDNPLRCDIC
						CVTFRTHRGLLRHNAL
22710	53078	A	22838	164	434	
22711	53079	A	22839	23	242	
22712	53080	Α	22840	187	325	
22713	53081	Α	22841	1	130	NCPMCWPGQSPRVPST*GRRNP
						PPKRRSVSAES*RHWPQSPSP
22714	53082	A	22842	614	774	QEPMRMCTMLANSSPSSSQATP
						PGL*SWNQPSLPARTLSRHEWT
						SCLGSSANL
22715	53083	A	22843	396	560	LFPHDGIKHKGKENP/PKGPPPP
						THPQTPPRPADAPLVATKSTLD
1						QAWALLAVATS
22716	53084	В	22844	124	205	
22716	53084	В	22844	124	205	<u></u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
		<u> </u>		<u> </u>		
22717	53085	A	22845	1	2355	MVGSRRISGVIPLPFTRRAGIFS
						DPRANMQPLWAPVPPATFVPTS
						LPPVPTARPPTSASCHCQQVPC
		1				AGLEVALPGNRTICWLSLWPS
						WQLDEDAQSPALESSSIRQFPT
						APRSGYPGTLCPSFLFGPRPRTC
						EYHSKPVLCAPSSRASPSSATTL
						STRTLWGPGAGSHPFGVHNTR
						LSPDLCPGKIVLRALKESGAGM
						PEQHKDPRVQENPDDQRTVPE
						VTGDARSAFWPLRDNGGPSPFV
						PRPGPLQTDLHAQSSEIRYNHTS
						QTSWTSSSTKRNAISSSYSSTGG
		l				LPGLKQRRGPASSRCQLTLSYS
						KTVSEDRPQAVSLGHTRCEKG
						ADTAPGQTIAPTASRPHINTSLH
						VEDKAISDCRPSRPSHTLSSLAT
						GASGGPPVSKAPTMDAQQDRP
						KSQDCLGLVAPLASAAEVPAT
						APVSGKKHRPPGPLFSSSDPLPA
						NSSHSRDSAQVTSMIPAPFTAA
						SRDAGMRRTRSAPAAAAAAPP
						PSTLNPTSGSLLN\EWMEAPHIS
		1				WPQPQLQHVPRGQ/NVRYNQR
		ļ				SQTSRTRS/CPQTKCQLQLPQLY
						GRPPGSKAEEG\PASSHCQLALS
ļ		l				SSNTVSEDGPQAVSSGHRCENK
						AGTAPGQTLAPRGGSPRSQASR
						PHINTALHVEDKAISDCRPSRPS
•		l				HTLSSLATGASGGPPVSKAPTM
			'			DAQQDRPKSQDSLGLLAPLASA
						AEVPSTAPVSGKKHRPPGPLFSS
22718	53086	A	22846	1	1200	MPEQDKDPRVQENPDDQRRVP
22/10	33080	<u> </u> ^	22040	1	1200	EVTGDARSAFRPLRDNGGLS/PP
		l				FVPGPGPLQTDLHTQR\LGYRV
		l				TVEDLDREKEAAFQRINSALQV
						EDKAISDCRLSRPSHTLSSLATG
j		İ				
						TSGLPAVSKAPSMDAQQERHK
-						SQDCLGLLAPLAPAAEVPSTAP
ļ						MSGKKHRPPGPLFSSSDPLPATS
						SHSQDSAQVTSLIPAPFPAASM
						DAGMRRTSPGTSAPAAAAAPPP
						STLNPTLGSLLEWIEALHISGPQ
						PQLQQVPRGQNQRSQTSRTSSC
						PKRK/CHLQLLQLYGRPPGTKA
						EEG\PASSHCQLTLSSSNTVSED
						GPQAVSSGHTHCEKTADTAPG
						QTLAPRGGFPRSQASRPCRHKF
						PLLPRRRGEPLMLPPPLELGYR
						VTAEDLDREKEVAFQRIKSALQ
		L				VEDKAI
L	L	Ь	<u> </u>	· · · · · · · · · · · · · · · · · · ·	L –	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22719	53087	A	22847	2	1762	YHSKPVLCAPSSRASPSSATTLS TRTLWGPGAGSHPFGVHNTRL SPDLCPGKIVLRALKESGAGMP EQHKDPRVQENPDDQRTVPEV TGDARSAFWPLRDNGGPSPFVP RPGPLQTDLHAQSSEIRYNHTS QTSWTSSSTKRNAISSSYSSTGG LPGLKQRRGPASSRCQLTLSYS KTVSEDRPQAVSLGHTRCEKG ADTAPGQTIAPTASRPHINTSLH VEDKAISDCRPSRPSHTLSSLAT GASGGPPVSKAPTMDAQQDRP KSQDCLGLVAPLASAAEVPAT APVSGKKHRPPGPLFSSSDPLPA NSSHSRDSAQVTSMIPAPFTAA SRDAGMRRTRSAPAAAAAAPP PSTLNPTSGSLLN\EWMEAPHIS WPQPQLQHVPRGQ/NVRYNQR SQTSRTRS/CPQTKCQLQLPQLY GRPPGSKAEEG\PASSHCQLALS SSNTVSEDGPQAVSSGHRCENK
						AGTAPGQTLAPRGGSPRSQASR PHINTALHVEDKAISDCRPSRPS HTLSSLATGASGGPPVSKAPTM DAQQDRPKSQDSLGLLAPLASA AEVPSTAPVSGKKHRPPGPLFSS SDPLPATSYHSRDTAQ

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SEQ ID NO:	,		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22720	53088	A	22848	292	2053	GSVAWSPFSPSPSSLCLWDLPV
	,					APTLWGPGAGSHPFGVHNTRL
						SPDLCPGKIVLRALKESGAGMP
						EQDKDLRVQENPDDQRRVPEV
						TGDARSA/FLAPAGQWSPLSLC
						AQSRASADIPPCPEVRNQI*PDIP
						DHLDELMHQPKCHLQLLQLCG
						RLAGAKVEEGASGAKEQGRDA
						*AGQGTPESQQNPDDQRTVPEV
						TGDARSTVRPLRDSGGLSPFVP
						RPGPLQTDLHAQSSEIRYNQTS
						QTSWTSSSTKRNAISSSYSSTGG
					ŀ	LLGLKQRRGPASSRCQLTLSYS
			ļ			KTVSEDRPQAVSSGHTRCEKA
						ADTAPGQTLAPRGGSPRSQGSR
						PRRRKIALLPHR*GELQVEDKAI
						LDCRPSWPSHTLSSLATGASGE
						PPVSKAPTMDAQQDRPKSQDC
						LGLVAPLASAAEIPSTAPVSGK
						KHRPPGPLFSSSDPLPATSSHSR
						DSAQVTSLIPAAFTAASMDVG
ŀ						MRRTRPGTSAPAAAAAAPPPST
						LNPTSGSLLN\EWMEALHISGPQ
			1			PQLQQVPRGQ/NVRYNQRSQTS
						RTRS/CPQTKCQLELPQLYGRPP
						GTKGCRHEKNVLCSKLFEGFGF
						IFVGFFFFAYMGILQLLII

SEQ ID NO:	SEQ ID NO: of peptide sequence	,	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22721	53089	A	22849	146	1789	RQPSTSGSSRSCPCARRSWWAR
						TARSGWWWELLGARRSPQTLH
				į		CPQATLITLPCPRPSSTTSGSAM
						T*RGPWRSPGCTTSFCPTSR/PV
						ERNIDQTDLHAQRSEIRYNQTS
						QISWTSSCTNRNAISSSYSSTGG
						LPGLKRRRGPASSHCQLTLSSS
					;	KTVNSDKFDANDPILKDQTQE
						WSGSATFTSDGKIRLFYTDYSG
						KHYGKQSLTTAQVNVSKSDDT
						LKINGVEDHKTIFDGDGKTYQN
						VQQFIDEGNYTSGDNHTLRDPH
						YVEDKGHKYLVFEANTGTENG
		l				YQGEESLFNKAYYGGGTNFFR
		1				KESQKLQQSAKKRDAELANGA
ļ		1	,			LGIIELNNDYTLKKVMKPLITR
		i				GSSRPSIREDENGRSQKPVHPEG
ł				, ·		DMTMNIKKIVNQATVLTFTTAL
1						LAGGATQAFAKENNQKAYKET
						YGVSHITRHDMLQIPKQQQNEK
1						YQVPQFDQSTIKNIESAKGLDV
						WDSWPLQNADGTVAEYNGYH
						VVFALAGSPKDADDTSIYMFY
		1				QKVGDNSIDSWKNAGRVFKDS
						DKFDANDPILKDQTQEWRGEA
22722	53090	A	22850	500	1034	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	Į.	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>				
22723	53091	Α	22851	1	2396	MDKYLLHGPPRPPPAIASRCLA
						RVPGCAYSSIMVALSLVPLCAH
					:	MPTPSGLEVALPGNRTICWLSL
						WPSWQLDEDAQSPALESSSIRQ
1						FPTAPRSGISPLCAPHSCLGQGR
						GPASHPTSTQTLDISGVKTTPRR
						FSVILQDLCSILGDSCLQFHLHR
						VPQQVSAVLLRVSGASPSPATT
						LSTRTLWGPGAGSHPFGVHNT
						RLSPDLCPGKIVLRALKESRAG
						MPEQDKDPRVQENPDDQRTVP
						EVTGDARSAFWPLRDNGGPSPF
						VPRPGPLQTDLHAQSSEIRYNH
						TSQTSWTSSNTKRNAISSSYSST
		ĺ				GGLPGLKQRRGPASSRCQLTLS
						YSKTVSEDRPQAVSSGHTRCEK
						GADTSPGQTIAPTGGSPRSQDS
						RPRRRKIPLLPRRRGEPLMLPPP
						LELGYRVTAEDLHLEKETAFQR
						INSALHVEDKAIPDCRPSQPSHT
						LSSLATGASGGPPVSKAHTMD
						AQQHRPKSQDCLGLLAPLASA
						AEVPSTAPVSGKKHRPPGPLFSS
		ĺ				· .
						SDPLPATSYHSRDSAQVTSLIPA
						PFTAASSDAGMRRTRPGTSAPA
						AAAAAPPPSTLNPTSGSLLN\EW
						MEAPHISWPQPQLQHVSRGQ/N
						VRYNQRSQTSRTRS/CPPRKCQL
						QLPQLYGRPPATKAEEG\PASSH
						CQLAHSSSNTVSEDGPQAVSSG
1						HTRCEKKAGAS/RWASRF*STH
						YGCTAGQTQVPRLPGPTGPPSi
22724	53092	Α	22852	1	3392	MDKYLLHDPPRPPPAIASRCLA
		ŀ				RVPGCAYSSIMVALSLVPLCAH
						MPTPSGLEVALPGNRTICWLSL
						WPSWQLDEDAQSPALESSSIRQ
ł						FPTAPRSGYPGTLCPSFLFGPRP
						RTCEYHSKPVLCAPSSRASPSSA
		İ				TTLSTRTLWGPGAGSHPFGVHN
						TRLSPDLCPGKIVLRALKESGA
						GMPEQDKDPRVQENPDDQRTV
						PEVTGDARSAFWPLRDNGGPSP
						FVPRPGPLQTDLHAQSSEIRYN
	1	1				HTSQTSWTSSST
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH
122,23		` `		"	"	RLFFLNGVFEHLGRAFVQRRKV
						DTEHQLLRAAVPANQEHRGLT
						QSQHVDGHSLLLLTLPNSFLCQI
						QEPSLG\SGSGALSGNVRIRPQA
1						Į -
00706	52004	 _ 	22067	12	220	QEAPEEPSVLTLLW
22726	53094	Α	22857	3	220	i

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22727	53095	A	22858	1	356	MAGPVKDREAFQRLNFVYQVS LRQGPHGDGARRPRVTAPLPQ AAHCVLAQDPENQALARFYL\H TERTIAKRLVLRRDPSVKEELL CRGCSSLLVPGLTCTPAPRDRC RGQRWTGTDLP
22728	53096	Α	22859	2	270	
22729	53097	A	22860	2	364	IFSALPAPRRKVGLNLAPVTEPR DQPWAMIIDVF\SRYSGSRGQPR QT\LT\KGELKVLMEKELPG\FL QSGKDKDAVDKL\LKDPGRPM GDAQGGTFSEVHPCSVAAIT\SA CHK\YFE\KAGLK
22730	53098	A	22861	282	469	DCGSELQAPPPPSLPGPLCPSIM PA/PPYASPPMMPMMGPPFRGM MPVGLAPGPMLLQYNTCS
22731	53099	A	22862	7	522	SVWWNSNGLQSNMPKFYCDY CDTY\LTHDSPSVRKTHCSGRK HKENVKDYYQKWMEEQAQSLI DKTTAAFQQGKIPPTPFSAPPPA GAMIPPPPSLPGPPRPGMMPAP HMGGPPMMPMMGPPPPGMMP VGPAPGMRPPMGGHMPMMPG PPMMRPPARPMMVPTRPGMTR PDR
22732	53100	A	22863	3	455	WFLIHDRTHTGEKPYKCNECV KAFHRSTCLHAHKRTHTGEKP YECNQ*GKAFNSSHSFQIHERT HTGEKPYECKECGKAFKGPSSV RRHERTHSTKKPYECKHCGKA LSYLTSFHNHLGMHTGEISHTC KICGKAFYSPSV\IQTHE*THT
22733	53101	Α	22864	1	1665	
22734	53102	A	22865	1	2650	

2988

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	-	
22735	53103	Α	22866	107	2384	MFGVHGTFLGIESRPPLLSLSPS
						LLYTCEMFQDPVACEDVAVNF
						TQEEWALLDISQRKLYREVML
	1					ETFRNLTSIGKKWSDQNIEYEY
					:	QNPRRSFRSLIEKKVNEIKDDSH
						CGETFTQVPDDRLNFQEKKASP
						EVKSCDSFVCAEVGIGNSSFNM
						SIRGDTGHKAYEYQEYGPKPY
						KCQQPKNKKAFRYRPSIRTQER
						DHTGEKPYACKECGKTFISHSG
						IRRHMVMHSGDGPYKCKFCGK
		l				AVHCLRLYLIHERTHTGEKPYE
	İ					CKQCGKSFTYSATHRIHERTHT
	ł					GEKPYECQCGKAFHSSSSFQA
	ļ					HKRTHTGGKPYECKQCGKSFS
						WCHSFQIHERTHTGEKPCECSK
			i			CNKAFRSYRSYLRHKRSHTGEK
						PYQCKECRKAFTYPSSLRRHER
		1				THSAKKPYE/CSFQYHERTHTG
	1					EKPYECKQCGKAFRCAPHLRR
						HGRTH/S/AEKPYECKECGKAFR
						SASHLQIHERTQTHIRIHSGERP
						YKCKTCGKGFYSPKSFQRHEKT
						HTAEKPYECKQCGKAFSSSSSF
						PYHERTHTGEKPYECKQCGKA
						FRSASILQMHAGTHPEEKPYEC
						KQCGKAFRSAPHLRIHGRTHTG
						EKPYECKECGKAFRSAKNLRIH
						ERTQTHVRMHSVERPYKCKIC
i i	1					GKGFYSAKSFQIHEKSYTGGKP
		l				YECGLQLPRTAAESVDSEKLQG
						PGKQETHVQLLSLYCLRRVLDC
22736	53104	Α	22867	ı	391	MKEHIERNPVTIKNVVKPSINT
		1				VPLEDMKAHTGEKPYERQKCE
						LNQRLRDYAFQYHESNSGVVS
						ADSETKSGHSGCLIKEMKNATK
						THNGITGGIIFNANHWRETNTIY
				·		TRIHI/GEKSCDYKKCGKAFNQ
						YSSLRRHES\HTGEKPYERQKC
					}	ELNQRLRDYAFQYHESNSGVV
	-					SADSETKSGHSGCLIKEMKNAT
						KTHNGITGGIIFNANHWRETNTI
						YTRIHINKQEKYQATHVSYAAN
22737	53105	С	22868	179	289	
22738	53106	A	22869	3	206	
22730	100100	l' `	122007	<u> </u>		

SEQ ID	SEQ ID NO:				1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	deterion, (-possible nucleotide insertion)
		1				
22739	53107	Α	22870	1	855	MPRAEPRATLGEQEKAGLPLG
						AWRLYLLRHFRKQTELRRSGSR
						DVTGALLVAAAVASEAVGSLR
1						VAEGGPNTLLLQVLR\SW\PWC
						NKELKTMEERKVKRRSPKSFSA
						HCTQVVNVQKKMSIPVSK\ST\G
						FSNPASQVNFTSRPKVKKEVM
						K\EKTKPQGLEEGQRHSSQLPIQ
						HSFLT\DVSRCSRKMERGLLSL\
						LNDFHSWKTFKHFGN\ECSI\EQ
						MEHVSGELQEKLARLNLELYG
						ELEELPEDKRKTASDSNLDRLL
						SDLEELNSSIQKLHLA\DAQDVP
22740	53108	IA	22871	67	598	SIRRHCGQRCLVRIRIWPLKLPP
		-				RLAVREGLRCGTLRDVMPPSM
						SGVVVGPPGP\PPGPCGDTTSGP
						APGTFGQASAPVSPPRRGHGEY
		l				LCSSPTPVFFPGLWQPPRAGFTA
					:	PERSWSCRGPRWQLRAAWAGR
						LSSSPVSALLTFKVIILSTGLLFG
						PLGGWKILSSLPEIFANLAT
22741	53109	A	22872	1	372	I BOO WILLBOOM BILLING IN
22742	53110	A	22873	li li	3961	FALAPEAGFYSLGQQPSTRISGS
						GCVHPVRVWPFSIEPAIGTLNV
		1				GESMQLEVEFEPQSVGDHSGRL
						IVCYDTGEKVFVSLYGAAIDMN
			}			IRLDKNSLTIEKTYISLANQRTIT
						IHNRSNIIAHFLWKVFATQQEE
						DREKYSKVTAVFSIAASSNSRF
			}			ALYESSYNSQRSPSEWACDDLI
		1				KEEKDETDEFFEECITDPLLREH
						LSVLSRTFANQRRLVQGDSKLF
						FNNVFTVEPLGWSERKEMAEIR
						DVOKKDTKV
22743	53111	A	22874	1248	1716	WYAEQSAHEATAQLSTPMAIR
				12.0		RAVVMPPEQRNRTKKEY/SLRI
						PFGLPGDRMCFFTHRNDD/PTG
						NEVTPPSLAADLGQQFGGDAT
						DRQPWMIHQMLIHQGSEQLCF
						VEHGINMLRLGEILHLAATRHE
						AHFLAFRQFFNHQRSCFALNLA
						RQDRITDF
22744	53112	A	22875	478	663	
22745	53112	A	22876	508	884	RQRHPDQRPGRSLFHGHGMCP
		['	TTPLQQQVDSQHNGDNNSESA
						P*QRRADSIHRSSLHIKKRPPSE
					İ	DVSRCFLCIDLAFRQGFTFNKK
						GIRVRIPFGLPGDRMCFFTHRN
						DDRPAMKLLPFAGNKV
L	1		L	L		DUNI AWIKEEI PAUINKY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
•	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22746	53114	Α	22877	1	506	AFCRFSILVGSAFISETDMCLLC
						PFPIGKAISVRLQYFKHETKQSI
						CLGNQKSPLVCPFFGEQGLDSS
						SSLDLPSSPNPAPTLCPCVQPPR
						GPVSVSVPVSVMGSRLAPLLSA
						*LFVSVILGRMVILKNPGVLGQ
		l				RQAGPSPGAPGLPSPSVRAPLG
						HKCSERSPSAT
22747	53115	A	22878	621	877	RRRRRRKQRRKQPWSSPARGA
						*PGPRRPSLRAAGTRRRKRRA
						SWTRCSNTCPGRCRTCSRSNQA
						SCPSLLHLLLLFGSFFSFY
22748	53116	Α	22879	1	1339	MPSYKGCGETQRHATNGTAAP
						PRRSSCRGDAIKLPHAEKKEGR
						PPASTSYPLRLGTRSARSSSHC
						WCTSWEDPPEATAGEEKRHPK
						SPAVCCGEGTVLTASTVTWRA
						EPARGEIWARLCPRDRPGSRAA
		ĺ				AVIWERPCCDRLIADRTGPPNSS
		l				EPHCRAATIPLPTWPLPEGQTLT
						LKALAKMAAPALLLLALLLPV
						GAWPGLPRRPCVHCCRPAWP/L
		1				WTLCPGEKGEAGVRGRAGRSG
				İ		KEGAA\GCPGPAGPQSARRGSV
						GPPGAACRRAYAAFSVGRREG
						LHSSDHFQAVPFDTELVNLDGA
		l				FDLAAGRFLCTVPGVYFLSLNV
]				HTWNYKETYLHIMLNR/QAA/V
						VLYAQPSERSVMQAQSLMLLL\
						RRATPVWSVPEELPAVLPGQDE
						LGSLQLHAHHPLADGLVRHEL
		ł				QPGLAGSSRTGHLVFRTDHGLR
	1					QHASHGCPVGPLHVPG
22749	53117	Α	22880	532	783	VGSTRGIVEGLIFILGEGGECYY
		Ì	İ			PRSLFTVANFTVGKRV\KTLNC
						PSTDKWINKMCHIHTVEYYSDI
		ŀ				KRNEIPMRATCRQTLKA
22750	53118	Α	22881	275	720	TPIHNCFKENKIPRNPTYKEHEG
		ł			<u> </u>	PLQGELQTTSQGNKRGYKQME
						EHSMLMG\GRINIMKMAILPKVI
		}				YRFTIPIKLPMTFFTELEKTTSKF
						IWNQKRARIAKSILSRKNKAG/
						AHHAT*LQTILQGYSNQNSMVL
		1				VPKQRYRPMEQSPQK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22751	53119	A	22882		609	MGRNQSRKAENSKNQSASSTP KDHSASPATEQSWMGNDFDEL TELGFRKSVMTNFSQLKEDVRT HHKEAKNLEKRLDKWLTRINSI EETLNDLMELKTMARELRDTC TSFSSRFDQVEEGEIQTTIREYY EHLYANKLENLEEMDKFLDTY TLPRLNQEEVESLNRPITGSEIE AIINSLPTKKIPGPDRFTA*RRP* MT*WS*KPWHENYVTHAQASV ADSIKWKKGKYK\HHQRIL*TP LRK*TRKSRRNG*IPGHIHSPKT KPGRS*IPE*TNNRF*N*GNN**P TNQKNSRARQIHSQILPEVQRG
22752	53120	A	22883	191	564	PNARKLRTLKKDHSPD**RRKL RKIK*MQ*KMIKGISQ\RSHRNT NYHQRTL*TPLHK*TGKSRRNG *IPGHIHPPKSKPGRSQIPEYTNN KI*N*GSN**PINQKKSRTGRIHS QILPEVQRGA
22753	53121	A	22884	337	589	TRRSRRNG*IPGHIHPPKTKPGR SRIPE*TNNRL*N*GSS**PTNQK KPRTRETDSRILPEVQRGAGTIP SETIPNNRKRGTPP
22754	53122	A	22885	162	539	TRKSRRNG*IPGHIHPSKTKPGR S*IPEETNNKF*N*GSN**PTNQK KPRTRQVHSRILPEPISHGMGKS WKQSL*KPAQDKAVPSLIAPIQ HIIGSSGQGNQARERNKGYSNR KRGSQIVSVCR
22755	53123	A	22886	329	459	NDDTFRKPHRLSPKSP*AADKQ LQQSLRIQNQCAKLRSIPIHH
22756	53124	A	22887	58	440	GLSRQFSPHSVNKAAGEVPTGR GPPQLSKANVARLPLSIPPLWA GHLWKKGGSPSQGIIDKTLISLG QST*GKGQLWVQLQRT*AFLPA GSEESSGSPSTA\PNSAKGQTPS SSGSLTRASSSSLYLW
22757	53125	A	22888	219	332	NEGKNLKGSQRERNTNYHQRT Q*TPLHK*TRKSRRNG
22758	53126	A	22889	1295	1435	QNQRQKPHDYLNRCRK/W/HFD KIQQPSMLKTLNKLGNDGTYL RIIRDI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	1	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22759	53127	Α	22890	2	648	YRFSGPTIRSIQNHTTHNQGRK
1				1		WHKKGIKKPRAQRYESLKGPE
						V*PSGQGVTCLLRADWLSPLLG
						SGTY\VGPKFLRNTRFAKKHNK
						KGLKKMQANN\AKAMSARAE
						AIKALVKPKEVKPKIPKGCS/RR
						KLRFDFAYIAHPKLWGRRCFVA
						RFCQGASGCSGPKGKAKVPKA
						KVKVS/PKAKAKAKAKDQTKA
						QAAAPALV\PAQAPKRTQAPTK
l						ASE
22760	53128	A	22891	3	370	QMYCVFNRNEDACRY/GS/AIG
						VLAFLASAFFLVVDAYFPQISN
						ATDRKYLVIGDLLFSG\VLASLA
						YQRYKAGVDDFIQNYVDPTPD
						PNTAYASYPGASVDNYQQPPFT
				1		QNAETTEGYQPPPVY
22761	53129	A	22892	11	297	(
22762	53130	A	22893	3	713	GGGSGGDGDMESGAYGAAKA
						GGSFNLRRFLTQPQV\VARAVV
						L\VFALIVFSCIYGEGYSNAHES
						KQMYCVFNRNEDACRYGSA\IG
						VLAFLA\SA\FFLVVDAYFPQISN
						ATDRKYL\VIGDLLF\SALW\TFL
						WFR/VGFCFLTNQWAVTK\PKT
						VLVGADSVRAAITFSFFSIFSWG
						VLASLG\YQRYKAGVDDFIQNY
						VDPTPDPNTAYASYPGASVDN
						YQQPPF\TQNAETTEGYQPPPV
22763	53131	A	22894	3	219	TQQTT (TQNAETTEGTQTTT V
22764	53132	A	22895	1445	1777	PVHLGSTCHQCRQKTIDTKTNC
22/04	33132	Δ.	22093	1775		RNPRLLGA/CRGQFCGPCLQNR
		1				YGEEVRDALLDPNWHCP\HCR
						GICNCSFCRQRDGRCATGVLVY
						LAKYHGFGNVHAYLKSLKQEF
						EMQA
22765	53133	1	22896	3	337	LIVIQA
22766	53134	A	22897	425	1209	TMSSTQFNKGPSYQLFAEIKNQ
22700	33134	^	22097	423	1209	LLFKCDPQKQAALCSWIEGLTR
						LSFSPDFQKSLKEGIVLCTLMN
						KLGSVPKI/NRLMQNWHQLENL
						SNFIKAMVGYDMNPMDLFEAN
						DLFESRNMTQNRIL/P/PLDHSTI
						SLQMGANKRSSQ/VGMTAPGTP
						QHIYNTKLGAHKCDNSSMSLH
						MGDTQSANHSVQGFGLGQQIY
						YPKYCPQ/GPAASGAPLDAGNC
						PGPGEAPEDPPCSQEEATTEAPE
						HAGSLHHRPVWVWGIFLYFRL
_				1		SFFFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22767	53135	Α	22898	156	559	QHIRETPGAQVHSCGKWNPQA
22707	33133	ļ .	22070	130		LWRTAWQLLT/DTKHTLGKPPS
						/MPHLVIYPLEISC\HTKTCTRML
						IASFFIITKTWKRPRCPSVDKWK
						NKPWCIHTGLLSSTVPPGEPGC
						PIPELSGLLMGLQTTSSRCSVRQ
						IVW
22768	53136	A	22899	161	359	
22769	53137	A	22900	151	383	
22770	53138	Α	22901	238	546	PQEHIMEHLEGVINKPRGRRCR
						PQELQLHYLQKCMDYDGNNLL
						DGFRTLPQPSTHV\HKEEGSEQ
						APLMSEDEL\INIIDGVLRDDDK
						NNDGY\IDYAEFAKSLQ
22771	53139	A	22902	1	819	MAAEDKLLLPQLPELFETSKQL
						LDDVEVVTEPAGSRIVQKKVFK
						GLDLLEKAAEMLSQL/DLVAEL
						ELPKTKNSAE\NHTANFSMAYP
						SLIAMASQRQAKRERHKQKEL
						EHRLSAIKSAVETGQADDERVR
						EYYLLHLQSLEEIESIDQEIKILR
						ERDSSREASTSNSSRQERPPVKP
						FILTRNMAQVKVFGAGYPSLAT
						MTVSDWYKQHWKYGALPDQG
						IAKATPEEFRKAAQQQEGQEEK
			į			EEEDDEQTLYRVREWDDWKGI
						HPRDYGSQQSMG
22772	53140	B	22903	238	410	
22773	53141	Α	22904	1	638	VAELELPKTKNSAE\NHTANFS
						MAYPSLIAMASQRQAKRERHK
						QKELEHRLSAIKSAVETGQADD
						ERVREYYLLHLQSLEEIESIDQE
						IKILRERDSSREASTSNSSRQERP
						PVKPFILTRNMAQVKVFGAGYP
						SLATMTVSDWYKQHWKYGAL
						PDQGIAKATPEEFRKAAQQQEG
						QEEKEEEDDEQTLYRVREWDD
						WKGIHPRDYGSQQSMG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22774	53142	A	22905	3	1083	RGNSRLRYSHEDELQLPRLPEL
	33112					F\ETGRQLLDEVEVAT\DPAGFR
						\IVQEK\VFKG\LDLLE\RAAEML
						SOSTLEGRECREWKDLSPTHLI
		}				TFWGP\AFQGA\LPLKQVNPSKR
						LRIILQR\AREHFI\NYLT\QCH\C
į						YHVGQSFEL\PKT\MNNSAENH
						TANSSMAYPMLGAM\ASQRQA
		l			ł	KIQRYKQKKELEHRLSAMKSA
		l				VESGQADDERVREYYLLHLQR
						W\VDISLEEIESIDQEIKILRERDS
						SREASTSNSSRQERPPAKPFILT
						RNMAQAKVFGAGYPSLP/TLW
		1				TVSDWYEQHRKYGSITGFRGIA
		ŀ				KSAPEGISGKAAQQQEEQEEKE
						EEDDEPNNFHRAREW\DDWKD
						TPSLGAYGNRQNMG
22775	53143	Α	22906	30	457	
22776	53144	Α	22907	401	1509	MWKMNCSLLQSKQMCNQLPP
						GSTRDGSWKFRQDPRLGKNA/C
						DCDSCIASEFGVRGYPTIKLLKG
						DLAYNYRGPRTKDDIIEFAHRV
1		İ				SGALIRPLPSQQMFEHMQKRHR
<u> </u>						VFFVYVGGESPLKYVTLKEMP
						AVLVFKDETYFVYDEYEDGDL
						SSWINRERFQNYLAMDGFLLYE
						LGDTDLNASSLRERKLILPEDIP
						LYVGKLVALAVIDEKNTSVEHT
						SELTVPTVVVLNTSNQQYFLLD
						RQIKNVEDMVQFINNILDGTVE
						AQGGDSILQRLKRIVFDAKSTIV
						SIFKSSPLMGCFLFGLPLGVISIM
						CYGIYTADTDGGYIEERYEVSK
						SENENQEQIEESKEQQEPSSGGS
22777	52145	-	22000	53	543	VVPTVQEPKDVLEKKKD
22777	53 145	A	22908	33	J343	LPISSLRTQTLFSSPLFPQHFEHY LALVSVLPFQLGRGRTAPAKKG
		1				GEK/RKGHSAINEVVIQEYTINI
						HKRIHGVRFKKRAPRALREIPK
						FAIKEVETLSVRIDSRFNKALCA
1						KGIRNVPYRIRVQLSRKRNEDE
						DSPNKRYTLVTYVPVSTFKNVQ
ŀ						TVHVHEN
22778	53146	A	22909	106	491	IGKGTNELMYGEK/RKGHSAIN
122,70	33170	[,				EVVIQEYTINIHKRIHGVRFKKR
						APRALREIPKFAIKEVETLSVRI
						DSRFNKALCAKGIRNVPYRIRV
						QLSRKRNEDEDSPNKRYTLVTY
						VPVSTFKNVQTVHVHEN
22779	53147	A	22910	1	321	
		1.,	1	<u></u>	<u> </u>	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22780	53148	Α	22911	313	402	R*GGEKKKGRSAINEVVTREYT INIHKRIHGVGFKKRAPRALKEI RKFAMKEMGTPDVRIDTRLNK AVWAKGIRNVPYRIRVRLSRKR NEDEDSPNKLYTLVTYVPVTTF KNLQTVNVDEN
22781	53149	Α	22912	1	3801	
22782	53150	A	22913	2	535	GGGMADTVLFEFLHTEMVAEL WAHDPDPGPGVSAGLRGEEAG ATKGQKMSLSVLEGMGFRVGQ ALGERLPRETLAFREELDVLKF LCKDLWVAVFQKQMDSLRTN HQGTYVL\QDNSFPLLPMGSG LQV/YLEEAPKFLAFTCGLLRSALYTLGIESVVTASVAALPVCKF QVVIPKS
22783	53151	A	22914	1	201	GDLPWEINPLSSCSLLHEKDPPT TSGPQT/DQPKKHHTNFKSGP\T GNRTVQLTWQPLPEPLELWPK AL
22784	53152	В	22915	226	522	
22785	53153	С	22916	1	705	
22786	53154	С	22917	1	462	
22787	53155	С	22918	1	505	
22788	53156	В	22919	1	1059	
22789	53157	С	22920	1	411	
22790	53158	A	22921	1	212	LTSFLHCCLSHHYHGAQIHPRH FLHLICRCHCHQYCHHAYLDPC PQDSHHLHWCWQPLPEPLELRP \RLSD
22791	53159	В	22922	1	864	
22792	53160	В	22923	1	635	
22793	53161	В	22924	1	874	
22794	53162	A	22925	1	566	MPSVRTGDPIILHAHTLNPAVW LAWAVEGEGSLTLNIWICPWQI EPLRHLAYTPGSDATSLTVPTR VISPLCVDSWLFTDPLMKTHAP ERTFLSFPAAAKRQAHLCVLAS NCDKPMYVKLVKAHCAEHQIN LIKGDDNKKLGEWEGPCTIDRE GKPCKVVGCSCAVVKDY/GKE SQAKDVIEQYFKCKK
22795	53163	A	22926	59	602	PTAMVEEGIAAGGVMDVNTAV QEVLKTALIHDGLARGIREAAK ALDKYVYQSQYCGFLQPEQNC HPREEGMEFMVLAQKF*MASV LPET*EK\RQAHLCVLASNCDEP MYVKLVEALCAEHQINLIKVD DNKKLGEWVGL\CKIDREGKPR KVVGCSCVVVKDYGKESQAK DVIEEYFKCKK

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22796	53164	С	22927	135	293	
22797	53165	В	22928	20	325	
22798	53166	Α	22929	677	1031	TPTMCLRSHPLPLALVSVILSFP
		l				QPSFSSSLINSSWSSPRSSLSSSSS
						RSPSSGRVNLHPPQTPSSH*SHL
						SSRSCCSAPLLTYHSSSLLLVCS
						VPFPFFPGFPGENPLFVLSGSAP
22799	53167	Α	22930	1	1011	MCLRVIFTIGAVSVILSFLSLFFII
			1			LDQLQLVIPQSSLSSSSNLEDPY
						SESYYAAARRPDSASTEPPPAG
					KSSAEYRAHFLAVSTEAERIRA	
				1	DLVLRARPLGPAHPRTLWPTPP	
				1	AGPFCRMVAHNQVA\ADNAVS	
						TAAEPRRRPEP/SPSSSSSSPAAT
						ARPRPCPAV\PAPAPGDTHFRTF
		ŀ				RSHADYRRITRASALLDA\LGF
		l	•			YWGPLSVHG\AHEPLRAEPVGT
		ŀ				FPVRDS\RQRNCFFALSVKMAS
						GPTSIRVHFQAGRFHLDG\SRES
						FDCLFELLEHYRGGSRADLLGA
	-					T\LRQRRVRPLQELCRQRIRRPP
						VG\REN\LARIPLNPVLRDY\LSS
	50160	<u> </u>		100	1166	FPFQI
22800	53168	В	22931	180	1466	
22801	53169	A	22932	100	207	CLOOCDCCNOLICCUDED A DAY
22802	53170	A	22933	408	3758	CLQQGPGGNQHCSHREEAPAV AACQEPPGHSEPESVLPLGAPP\
						SEEDGRQQVGSSRLRHIMAEMI
						ATEREYIRCLGYVIDNYFPEME
						RMDLPQGLRGKHHVIFGNLEK
						LHDFHQQHFLRELERCQHCPLA
		ŀ				VGRSFLRHLIGQKQFPESPSGER
		İ				PGLTSSHAPCPWMPHYLPEGPA
		ŀ				CPVLRIPCEAFNWKHVECTDPV
		ļ				VSPPKQSPLLFQEEQFGMYVIY
		1				SKNKPQSDALLSSHGNAFFKVI
		1				PLGPPPTACPALS
22803	53171	A	22934	3	355	EAAELDAMCTATRQWSRAGQ
22003	331,1	ľ.				GVTGRPCQGRLPRCGLRSTITP
						KPIPGGHRKKPLLGAL\GKHTLS
		1				YVERWHWEGHSGI*TRDRQNF
						ERRLTVPPSRQPFAPTDETRCG
						DDRISFCNYV

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22804	53172	Α	22935	1928	5312	CLQQGPGGNQHCSHREEAPAV
						AACQEPPGHSEPESVLPLGAPP\
						SEEDGRQQVGSSRLRHIMAEMI
						ATEREYIRCLGYVIDNYFPEME
						RMDLPQGLRGKHHVIFGNLEK
						LHDFHQQHFLRELERCQHCPLA
			,			VGRSFLRHLIGQKQFPESPSGER
						PGLTSSHAPCPWMPHYLPEGPA
						CPVLRIPCEAFNWKHVECTDPV
						VSPPKQSPLLFQEEQFGMYVIY
						SKNKPQSDALLSSHGNAFFKDK
						QRELGDKMDLASY
22805	53173	Α	22936	3	306	
22806	53174	Α	22937	3	460	LQRQRQHPAAAPAVPVRCFTFC
						FTDIVIMPKRKSPENTEGKDGS
						KVTKQEPTRRSARLSAKPAPPK
						PEPKPRKTSAKKEPGAKISRGA
						KGKKEEKQEAGKEGTAPSENG
						ET\KAEEIHISRSTVNVSTSRGTP
						PSTLSVKGQIETVRVKGTEN
22807	53175	A	22938	3	285	SSCPAREQLQRQRQHPAAAPAV
22007	23 1 / 2	-				PVRCFTFCFTDIVIMPKRKSPEN
						TEGKDGSKVTKQEPTRRSARLS
						/AVFV*FKLSLQKPAPPKPEPKP
						RKTSA
22808	53176	A	22939	1	413	
22809	53177	Α	22940	2	454	AEVQVEEAGLRRALRVSSCSGR
						GSIQRRFYAVPVRCFTFCFTDIV
						IMPKRKSPENTEGIDGS\KVTKQ
						EPTRRSARLSAKPAPPNPEPQPR
						KTSAKKEPG\AKISRGAKGKKE
						EKQEAGK\EGTAPAENDETKAE
1						EAQKTESVDNEGRMNCP
22810	53178	В	22941	55	186	
22811	53179	Α	22942	2	397	
22812	53180	В	22943	339	470	
22813	53181	Α	22944	1	139	
22814	53182	Α	22945	1	416	MPLLYHLSRCAYVQQCNFANP
						VVSALCTCLVKPVVEKCTPYA
						VLLEALALRNVRQLEDLVIEAV
						YADVLRGSMDQRNQRLEVDYS
						IGRDIQRQDLSAIARTLQEWG/C
					}	HQHLLSLRSAASQRQQCGYHK
		L				LHDKQRLLRG
22815	53183	Α	22946	315	570	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>				
22816	53184	Α	22947	933	1841	RPAEPDPGSGHPEPKSPTLSAV
						MSAEVKVTGRNQEQFLLLAKS
						A\KGAVLATLIHQVLE\APGVY
						VFGELLDMPNVRELAESDFAST
						FRLLTVFAYGTYADYLAEARN
						LPPLTEA\QKNKLSDTFSVV\TL
						AAKVKCIPYAVLLEALAL\R\NV
						RQLEDLVIEAVYADVLRGSLDQ
						RNKRL\EV\DYSIGRDIQRQDLS
						AIARTLQEWCVGCEVVLSGIEE
						QVSRANQHKEQQL\GLKHQIES
						EVANLKKTIKVTTAAA\AAATS
						QDPEQHLTELREPAPG\TNQRQ
			,			P\SKKASKGKGLRGSAKIW\SKS
22817	53185	Α	22948	1	1319	, and and to real thin it to the
22818	53186	A	22949	365	1331	RAGGGAPWRFTGTFATVAPET
22010	33100	` `	[22) 1)			AADLSLHRHLLSRTTSAWGTA
						CITVPIEPPTWPTWAVPIQWGG
						SATCIQEDKLRQGILYGYSQQV
			1			NQAMDSGLRP\VVDSVNAIRVP
						QDYVTQSVPLREMNGSLGVLA
						1 3
	}		!			QQLQNAKLQADAAHSALKQSD
						DLKPVFDQAFTKVVTTPADAL
						QPLIPAAQTFTQQLVMVGDYIA
						QQGTQECGKTFSRASYLVQHS
]					RIHIGKKPYECKECGKAFSSGS
						YLVQHQRIHTGEKPYECNKCG
						KAFTVYGQLIGHQSVHTGEKPF
						ECKECGKAFRLNSFLTEHQRVH
		<u> </u>				TGEKPFKCKKCGKTFRYSSAP
22819	53187	A	22950	2	206	GGGQGATSFGRPESGFIRHQPPP
						PCRRSSTPTRSKLYTL\SAPE*RS
	<u> </u>	<u> </u>				VPTSAPGPQDRPPGSVSKKSW
22820	53188	A	22951	571	1102	MGSVSLGNPQGPFQFTRLPLAN
						VIHPTFFWETGPRGADSLGAKG
		I				RKWQPTFTSRCNLKVYLGRAG
						SQAATGGPSPSLPNLHVKKLRP
			ļ			RKAEAWPGRRNTGKVSGR/LA
	ļ	1				ALRAPYGESFARAAGPRDAPS/
						QQPRPQPEHAPTLISLGSNFGG
						MVEESWCRMNPDSGRPKEVAP
	1					WPPPSTK
22821	53189	Α	22952	1	591	MGPHWKDFKEIDVKKYIDALSI
						HTYTHTHTHADITDGDHGG
						GALLELEINSHLKVQFRELNITA
						AKETEDGGGGKAIIIFVPFPQLK
				[SFQKIQVWLRRILPKPTRKSCT\
						QNMHWPPRSRTLKAMRDSIPE
						DLVF\PSEIMCMRIHVKLDGSRL
						IKAHLDKAQQYHVERNAETFS
[GVYMKLTGKDINVEFPEFQL
L	<u> </u>		L	l	l	O V TWIND TONDIN VERTERQL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22822	53190	Α	22953	2	508	APPFELPRAAGIRHEGKFSQEK
						AMFSSSAKIVKPNGEKPDEFES
		l				GISQALLELEMNSDLKAQLREL
]	NITAAKEIEVGGGRKAHIFVPVP
İ		Ì				QLKSFQKIQVRLVRELEKKFSG
						KHVVFIAHRRILPKPTRKSRTK
		1				NKQK\RPRSRTLTAVHDAIPEDL
						VFPSEIVGKRNR
22823	53191	Α	22954	145	395	
22824	53192	A	22955	536	717	
22825	53193	Α	22956	1	729	
22826	53194	Α	22957]	1158	
22827	53195	Α	22958	1	2869	MAHLRSCCSSLYQPGHLLPPHN
						PIQDHPLTAPSLTHLTRRAYFTD
						EKTKGGQSAYAVSLLRECVKL
ļ						RPSDPTVPLMAAKVCIGSLRW
						MAAKHLAGVLLHSLSEECYWS
						PLSHPLPEFMGKEESSFATQAL
						RKPHLYEGDNLYCPKDNIEEAL
						LLLLISESMLANKLLLLEMERV
						QSPGVESGVDRQLSPYSDTNVN
						AQNAQIVPVAAKSAQNEVPGY
						PEETEKSRVQSNLLTQFQDLRL
						LLAAPECLPPRMSPPS
22828	53196	В	22959	28	139	
22829	53197	Α	22960	60	812	LASPFPVPLHRCSAGSQPPGPVP
						EGLIRIYSMRFCPYSHRTRLVLK
						AKDIRHEVVNINLRNKPEWYY
						TKHPFG\HIPVLKTSKCQLIYES
ł						VIACEYLDDAYP\GRKLFPYDP
						YERVRQKMLLELFGKVPHLPK
						EFPGSVEDVGENAPNLKAALR\
						QEFSNLEEILEYQNTTFFG\GTCI
						SMIDYLPL/WPWFER\L\ECMGY
						WNCVSHT\PALRLWISAMKWD
						PTVC\ALLMDKEHF\QGFLNPYF
22820	52100	_	22061	2	251	QNNPNAFDFGLC
22830	53198	Α	22961	2	251	ILLDFKPWDDETDMAQLEACV
						CSIQLEELV*GASKLVPMGYGI
]					RKLQIQCVVEDDKVGTDLLEEI
	l	l				TKFEEHVQSVDIAAFNKI

3000

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
į	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22831	53199	Α	22962	353	1071	APLWPRAERKFYKQMNGPVAS
						ASRQSSAPG/NDHSELVLQIASL
						EVENQSLRGVVQELQQAVSNL
			ŀ			EAWLN\IRRREKSSPGH/RSTVP
						QTQHVSPMRQVERPSPRPPYYD
<u> </u>						PGQEAEDNEDDDIDLDNKEED
						KEARRLREERLRLYAKKAKNP
į						SLVAKSSILLDIKPWDNKTDMT
						QLEACVRSIQLDGLVSGASKLA
	•					SVGYGIRKMQIQCVVEDDKVG
						TDLLEEEITKFEEHVQSVDIAAL
22832	53200	Α	22963	3	482	PWANEAPAKK/PATPAEG/DAR
						DDDIDLF\GSDNEE\EDKQAGHS
ŀ						CGEER\LR\HYAGEEGPRKPAL/
						VWPKSLPSLLDVK\PWDDDDGP
						WPSLEACVR\SIQLDGLVWGGF
i						PSLVPVG\YGIRK\LQ\IQCVV\ED
ļ						DKVGTDFPWRREIT\KFEEARA
						RSVDIASFSTRS
22833	53201	Α	22964	1	2835	
22834	53202	В	22965	571	1337	
22835	53203	Α	22966	1	489	
22836	53204	Α	22967	1	618	
22837	53205	Α	22968	3	5227	SWSRVRPRTRPEGLRQLKMRT
						GWATPRRPAGLLMLLFWFFDL
						AEPSGRAVN\DPLTIFHGNTGKC
						IKPGYGWIVADHCDETEDKLW
ŀ						KWVPQHRLFHLHSQKCLGLDIT
						KSVNELRMFSCDSSAMLWWKC
						EHHSLYGAARYRLALKDGHGT
						AISNASDVWKKGGSEESLCDQP
						YHEIYTRDGNSYGRPCEFPFLID
						GTWHHDCILDEDHNGPWCATT
						LNYEYDRKWGICLKPENGCED
						NWEKNEQFGRCYQFNTQT
22838	53206	Α	22969	3	453	GQSPILVIYHETKRPSGISERFSG
			j			STSGNTATLTITGTQAMDEADY
						YCQAGRDFGGGTKLTVLGQPK
						AAPSVTLFPPSSEELQANKATL
1						VCLISDFYPGAVTVAWKAHSN
						PVKAGVETTTPSKQSNNKYAA
L						SSYLSLTPEQ*KSHRSYSC
22839	53207	A	22970	2	354	
22840	53208	Α	22971	1	315	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22841	53209	A	22972		712	MAWTLLFLTLLTQGTGSWAQS ALTQPASVSGSPGQSITISCIGTS SDIGAYYFVSWYQQYPGKAPK LLIFDVSGRPSGISSRFSGSKSGN TASLTISGLQAEDEADYYCASY ARSSTVVFGGGTSVTVLGQPKA APSVTL\FPPSSEELQANKATLV CLISDFYPGAVKVAWKADGSP VNAGVETTTPSKQSNNKYAAS S\YYLSLTPEQWKSHRSYSCQV THEGSTVEKTVAPTECS
22842	53210	Α	22973	1	1779	
22843	53211	Α	22975	1	2001	
22844	53212	A	22976	292	1497	TGQLNGFHEAFIEEGTFLFTSES VGEGHP\DKI\CDQIS\DAVLDA HLQQDPDAKVA\CETVAKTGMI LL\AGEITSRAAV\DLPRKVVRE SCLNTLEYDDSSKGF*LQRLVN VLVALEQQSPDIAQGVHLDRNE EDIGAGDQGLMFGYATDETEE CMPLTIVLAHKLNAKLAELRRN GTLPWLRPDSKTQVTVQYMQD RGAVLPIRVHTIVISVQHDEEVC LDEMRDALKEKVIKAVVPAKY LDEDTIYHLQPSG/RTFVIGGPQ GDAGLT\GRKII\VDTYGGWGA HGG\GAFSGKDYTKVDRSAAY AARWVAKSLVKGGLCRRVLV QVSYAIGVSHPLSISIFHYGTSQ KSERELL*IVKKNFDLRPGVIVR DLDLKKPIYQRTAAYGHFGRDS FPWEVPKKLIY

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)		
22845	53213	A	22977	67	1579	FFVRSLSHTDTNMNGQLNGFH		
	,					EALIEEGTFLFTSKADGEGHPD		
						KICDQISEAVLDAHLHQDPDAK		
						VACETVAKTGMILLAGEITSRA		
						AVDYQKVVREAVKHIGYDDSS		
						KGFD\YKTCNVLVALEQQSPDI		
						AQGVHLDRNEEDIGAGDQGLM		
				,	ļ	FGYATDETEECMPLTIVLAHKL		
						NAKLAELRRNGTLPWLRPDSK		
						TQVTVQYMQDRGAVLPIRVHTI		
						VISVQHDEEVCLDEMRDALKE		
						KVIKAVVPAKYLDEDTIYHLQP		
						SG\RFVIGGPQGDAGLTGRIIIVD		
						PYDGWGAHEEGAFSGKDYTKV		
						\SRSAAYGARWVARSLVKGGL		
						CRRVLVQVPGPPKKPDEDLYLE		
					[CEPDPVLALTQTLSFQVLMPSG		
						PLPRTSVVPREQQMLPLKDSDL		
						LTQPWYSGNCDRYAVESALLH		
						LQKDGAYTVRPSSGPHGSQPFT		
						LAVLLRGRVFNIPIRRLDGGRH		
						YALGREGRNREELFSSVAAMV		
						QHFMWHPLPLVDRHSGSRELT		
22846	53214	A	22978	3	325			
22847	53215	A	22979 22980	15 7	1014 592	LLCSGRHSLQGVMGILSFLPVL		
22848	53216	A 22980	22900	1	392	ATESDWADCKSPQPWGHMLL		
						WTAVLSLAPVAGTPAAPPKAV		
						LKLEPQWINVLQEDSVTLTCRG		
						THSPESDSIQWFHNGNLIPTHTQ		
						PSYRFKANNNDSGEYTCQTGQ		
						TSLSDPVHLTVLSGQWRKAPG		
						WTWEGQDG*NLLSGRGLQERG		
						VACLLGSIAVSCLSTYQWLFLP		
22849	53217	A	22981	32	1150	EKAVTAVLWAPARLQGVMGIL		
		-				SFLPVLATESDWADCKSPQPW		
				1		GHM\LLW\TAVLFLAPVAGTPA		
						APPKAVLKL\EPQW\INGGSRED		
						SVTL\TC\RGTHSP\ESDSIQWFH		
						NG\NL\IPTHTQPSLQVSRANNN\		
						DSGE\YTCQTGQTSLSDPVHL\T		
						VLSEWLVLQTPHLEFQEGET\IV		
						LRCHS\WRDKP\LVKVTFF\HNG		
						KSKKFSRSDPNFSIPQANHSHSG		
						DYH\CTG\NIG\YTLLFIPRLLTIT		
						VQAPSSSPMGIIVAVVTGTAVA		
						AIVAAVVALIYCRKKRISANPT		
						NPDEADKVG\AE\NTITYFTSQW		
						HPGWLWEEPDDQNRILVSIVLA		
İ								-
1	i .	1	1	Į.		LGIWRRNSEREDLVFPGLKFPL		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22850	53218	A	22982	1	471	MKSSFQGHNKEKREKKGPSLFS
22630	33210		22902	['	7 / 1	AAYGGGSCLLLSIMAALRPLVK
						PKIVKKRTKEFIWHQSDQHVKI
						KCNWRKPRG\QILMPNTGHGSN
						/KKTKHMLPSGFRKFLVPSVKE
						LEVLMCNKSPCAEIAHSVSSK\N
						KTIVERAAQ/LAVRVTNPNARL
						RSEENE
22851	53219	A	22983	103	564	GIFRGIMAALRPLVKPKIVKKR
22831	33219	A	22983	103	304	i e
						TKKFIRHQSDRYVKIKRNWRKP
						RGIDNRVRRRFKGQI\LMPNIGY
						G\SNKKTK\HMLPKWLSGKFLG
		1				SHQRSREL/EKLLA*CCNKFFTC
						CPRFVSHCFSFQGTGKGLRGER
22252	50000	 	22004		160	SLPNWAFRVTNPNARLRSEENE
22852	53220	A	22984	3	177	DEACACA CALIDEAD A DEPOSICA A DE
22853	53221	A	22985	74	735	RFAGAGAIPEARARPPDVQAAE
						EEKEMDLPDSASRVF/CGDASS
						NMGEHRMNVNAIIL\AQKNML
						D\RFEKT\NEMLLNFNNLSSARL
		1				ASRLSERFLHHTRTLVEMKRDL
		İ				\DSIFRRIRTLKGKLARQHPEAF
						SHIPEASFLEEED\EDPIPTQAPT
						TTIATSE\QSTG\SC\DTSPDTV\S
ĺ						PSLEPLASEDLSHVQALASPAI\
						NGPQPDQIDEEMTGRIALPARC
22854	53222	Α	22986	3	275	MRTVAEKVDAVRIVHIHSVI\IL
						RRSDKRKDRVEISPEQLSAAST
						EAERLAELTGRPMRVVGWYHS
						HPHITVWPSHVGKYHGVAMFI
		ļ				VFRN
22855	53223	Α	22987	1	886	AKMAVQVVQAVQAVHLESDA
						FLVCLNHALSTEKEEVMGLCIG
						ELNDDTRSDSKFAYTGTEMRT
						VAEKVDAVRIVHIHSVIILRRSD
						KRKDRVEISPEQLSAASTEAER
						LAELTGRPMRVVGWYHSHPHI
						TVWPSHVDVRTQAMYQMMDQ
						GFVGLIFSCFIEDKNTKTGRVLY
						TCFQSIQAQKSSEYERIEIP\IHIV
						PHVTIGKVCLESAVELPKILCP/
						QEEQDAYRRIHSLTHLDSVTKI
						HNGSVFTKNLCSQMSAVSGPLL
						QWLEDRLEQNQQHLQEL\HQE
						KEELMQELSSLE

SEQ ID	1	1	SEQ ID NO:		I .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hed	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		097340,217	sequence	or peptide sequence	deteriors, (-possible nacreotide insertion)
22856	53224	Α	22988	10	988	LKAMSWAKMAVQVVQAVQA
						VHLESDAFLVCLNHALSTEKEE
ŀ		l		<u> </u>		VMGLCIGELNDDTSRSDSKFAY
		ĺ				TGTEMRTVAEKVDAVRIVHIHS
		ŀ	1			VIIL\RRSDKRKDQVEISPGQLSA
						ASTEAEKLAELTGHPMRVVGW
						YHSHPHITVWPSHVDVRTQAM
		ļ				YQMMDQGFVGLIFSCFIEDKNT
		ŀ				RAGRVLYTCFQSIQAQKSSESL
						HGPQDFWSSSQHISIEGQKEEER
						YERIEIPIHIVPHVTIGKACLESA
						VELPKILCQEEQDAYWRIHSLT
						HLDSVTKIHNGSVFTKNLCSQM
						SAVSGPLLQWLEDRLEQNQQH
1						LQELQQEKEELMQELSSLE
22857	53225	Α	22989	1	1362	
22858	53226	Α	22990	284	542	KLFQKKNKKSSCISDNFFFLLAI
						MVNLLQIVRDHW\VH\VLVPM
						GFVIGCYLDRKSDERLTAFRNK
						SMLFKSFPHRELQPSEEVTWK
22859	53227	Α	22991	272	1241	FPGGAACDICWVKEENGSLMS
						MKMGWKAKSCLPVTVHPRCL
						TPYSARLSSTFPL*NSITTGP*QS
						PACKRPF*LTTC*GGSRRNSNR
						KAA*GPCSPPPPSPPPSPAT\LPR
		l				GPAGLQPPGVPVLPPLRPRKHY
						APGGLP\PPASLLEDDDDTFCTS
						QAMQPTAPTKLSPPALLPEKDS
						FSSALDEIEELCPTSTSTEAATA
						ATDSVKGTSSEAGTQKLDGPQE
		1				SRADDSKLMDSLPGNFEITTST
						GFLTDLTLDDILFADIDTSMYDF
						DPCTSSSGTASKMAPVSADDLL
						KTLAPYSSQPVTPSQPFKMDLT ELDHIMEVLVGS
22860	52220	A	22992	1	3234	MELKCVCLKNRSPSAEEQPSLL
22860	53228	A	22992	1	3234	PEPSAQESHGGHPRSALYGRER
						LTGOPPSSPHPNPAPWASCPAVI
						DPVPIYSSVAEAYDSGTEGIPTV
						PGLR\HVESGSLRATAQRLPPER
		1				RV/QQRTTAPPPRQPPRRPLRAS
						LSAAPPEPREPVVAAGPEPGSL
						QRPARTALPNRPEPQAASPGTL
						VEKYAAPVATQDLLDENPCTS
1						RLGVVDLQAVLTLRKLENLRS
						GAQTPLEGGQSGRVGPHALRW
		1				NRGEEGERLLC
22861	53229	A	22993	307	1906	
22862	53230	A	22994	1	1752	
		I			1	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22864	53232	A	22996	3219	3407	DQVVWTKRLQDAILVLV*EFRL HSPALRLCVMKQVGMSQGELE WGQFLKLSSRNRKRSGERI
22865	53233	Α	22997	1	588	
22866	53234	Α	22998	1	917	
22867	53235	Α	22999	1	1692	
22868	53237	A	23001	68	636	MKRIRDYYERSHANNLDNLEE MDELLETHNLPGLNHEETENLN KPITTSLSPLAGVSSTKKNTKNE LSIVLPFTGSILPVGFATRPRGL ASSNPILLHMVAFLFANGTEEL PVPLMTPTRQKKVIPLIPLMVG LGVSASTIALGTGITGISTSVTTF RSLSNDFSASITDISQTLSVLQA LVDSSAAVVLQGQLRGLDLLT AEKGGLCIFLNEECCFYLNQSG LVYDNIKKLKDRAQKLANQAR QNLDSMLHGTGMKSDSDQKKS ENGVTLAPEDTLPFLKCYCSGH CPDDAINNTCITNGHCFAIIEED DQGETTLASGCMKYEGSDFQC KDSPKAHVRRTIECCRTNLCNQ YLQPTLPPVVIGPFFDGSIRWLV LLISMAVCIIA\MIISSSCFCYKH YCKSISSRRRYNRDLEQDEAFIP VGESLKDLIDQSQSSGSGSGLPL LVQRTIAKQIQMVRQVGKGRY GEVWMGKWRGEKVAVKVFFT TEEASWFRETEIYQTVLMRHEN ILGFIAADIKGTGSWTQLYLITD YHENGSLYDFLKCATLDTRALL KLAYSAACGLCHLHTEIYGTQG KPAIAHRDLKSKNILIKKNGSCC IADLGLAVKFN\SDTNEVDVP\L NTRVGTK\RYMAPEVLDESLNK NHFQPYIMADIYSFGLIIWEMA RRCITGGIVEEYQLPYYNMVPS DPSYEDMREVVCVKRLRPIVSN GVTSNPKALLGALIAPAFASLPE GRGKRIQALNMFRPKNSWRLG LKYPPSSKEKTQVSAHKMPLSPG LLLLLFSGATATA/APLPLEGGP
22870	53238	A	23002	1	144	TG\RDSEHMQEAAGIRKSSLLTF LAWWFEWTS\QASAGPLIGKEA REVARRQEGAPP\QQSARRDRM PCRNFFWKTFSSCK GIRLPSSADWQKTEAQKRREQL
						\LLDELVALVNKRDALVRDLDA QEKQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22871	53239	Α	23003	2	667	RWNSSDPDADRTTLNHADHSS
						KIVQHRLLSRQEELKERARVLL
						EQARRDAALKAGNKHNTNTAT
						PFCNRQLSDQQDEERRRQLRER
]		ARQLIAEARSGVKMSELPP\YG
				1		EMAAEKLKERSKASGEQNSKL
						VDLKLKKLLEVQPQVANSPSSA
:				1		AQKAVTESSEQDMKSGTEDLR
						TERLQKTTERFRNPVVFSKDST
						VRKTQLQSFSQYIENRPEMKRQ
						RSIQED
22872	53240	Α	23004	1	1437	
22873	53241	Α	23005	1	2009	MAKPYLYKKYGNQQGVAAIPD
}						VPAAWEAEPSPIPSPVLGRKPN
ł		ŀ				ASQSLLVWCKEVTKNYRGVKI
						TNFTTSWRNGLSFCAILHHFRP
						DLIDYKSLNPQDIKENNKKAYD
						GFASIGISRLLEPSDMVLLAIPD
						KLTVMTYLYQIRAHFSGQELN
		}				VVQIEENSSKSTYKVGNYETDT
	[İ				NSSVDQEKFYAELSDLKREPEL
						QQPISGAVDFLSQDDSVFVNDS
						GVGESESEHQTPDDHLSPSTAS
						PYCRRTKSDTEPQKSQQSSGRT
				İ		SGSDDPGICSNTDSTQAQVLLG
						KKRLLKAETLELSDLYVSDKK
						KDMSPPFICEETDEQKLQTLDIG
						SNLEKEKLENSRSLECRSDPESP
						IKKTSLSPTSKLGYSYSRDLDLA
						KKKHASLRQTESDPDADRTTL
						NHADH\SSKIVQQDEERRRQLR
						ERARQLIAEARSGVKMSELPSY
						GEMAAEKLKERSKASGDENDN
				}		IEIDTNEEIPEGFVVGGGDELTN
		ŀ				LENDLDTPEQNSKLVDLKLKKL
						LEVQPQVANSPSSAAQKAVTES
		ŀ				SEQDMKSGTEDLRTERLQKTTE
						RFRNPVVFSKDSTVRKTQLQSF
						SQYIENRPEMKRQRSIQEDTKK
						GNEEKAAITETQRKPSEDEKGF
						KDTSQYVVGELAALENEQKQI
						DTRAALVEKRLRYLMDTGQNL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22874	53242	A	23006	145		LDEYEARLTLANLDDFEEDNED DDENRVNQEEKAAKITELINKL NFLDEAEKDLATVNSNPFDDPD AAELNPFGDPDSEEPITETASPR\ KTEDSFYNNSYNPFKE\VQTPQ YLNPFDEPEAFVTIKDSPPQSTK RKNIRPVDMSKYLYADSSKTEE EELDESNPFYEPKSTPPNNLVN PVQELETERRVKRKAPAPPVLS PKTGVLNENTVSAGKDLSTSPK PSPIPSPVLGRKPNASQSLLVWC KEVTKNYR
22875	53243	A	23007	1	165	RPAAPRHERRPAADHGHHRPEP GRGTGQKACSELPSDEACS/GIA IG*SLLCSACSVRSRKRQW
22876	53244	В	23008	26	625	
22877	53245	Α	23009	325	526	
22878	53246	A	23010	1946	1479	LLQWRHGLLNPINSPYNSPILPV QKPDKPYRLVQHLCLIYQIVLPI HPMVPNPYTLLSSIPFTTHYSVL DLKHAFFTIPLYPSSQPLFAFTW TDPDTHQAQQITLAVLPQCFIDS PHYFSQAQISSSSVTYLGIILMK THIGLGAVEQGVVLVGEARAA QEPMEWVGGSGMVGCRSRALP RGKAAKARREIERSAVTIVPVL DFNPAFHIIPDTTPDHHDCISLIH LTFTPFPHISFFPVPHLEHTWFID GSSTR
						QGELCESFDGKSLDGRDERVVI CFDAAAGLHRACSIRGEQQNL GTHYFTWQRLFAEVIKLGGLM KPKARNASSLWKEKTKNGFFS LETPEERQTSDLHKYQLAFRLF FIKYKNPAQFMAHLATTLRHFT ALDPESSHNFQNLFSSSHLMHIL SAPRLLRLYSRFVESPTVTIVPG PDFNPASNIIPDTTPDPHDCISLI HLAFTPFPYISFFPVPHPDHTWF IDGSSTRPNCHTPAKAGYAIVSS TSIIEATALPPSTTSRQAELIALT QALTLAKGLRINIYTD/SHILHH HAVLWA/E/RGFLTMQGSSIINA SLIKTLLQAALLPKEAEVIHCKG HQKASDPITQGNAYADKVPSL AHAPTLTLAPVLQCPLREVAGT EDDNGKAVLEIPSNIRKSAHVP VQTWKGWDPTNLLGGWFSNL GGFKTLVGTVIFIIGFLLFLPCVI PRIIKAIKTLVETTVSRQIIQTILL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22880	53248	A	23012	854	1777	
22881	53249	A	23013	27	496	
22882	53250	Α	23014	117	210	
22883	53251	A	23015	124	414	
22884	53252	A	23016	3	627	
22885	53253	Α	23017	l	567	
22886	53254	A	23018	44	551	GGGRRSAPLQLGFMSRQAKDD FLRHYTVSDPRTHPKGYTEYKV TAQFISKKDPEDVKEVVVWKR YSD\FRK\LHGD\LAYTHRNLF\R RLEEFPAFP\RAQ\VFGRF\EASV DSRER\RKGAEDLLRFTVHIPAL NNSPQLKEFFRGGEVTRP\LEVS RDLHILPPPLIPTPPPG
22887	53255	Α	23019	25	488	
22888	53256	С	23020	9	119	
22889	53257	A	23021	52	241	
22890	53258	В	23022	4	402	
22891	53259	A	23023	249	984	GSTMYKMNICNKPSNKTAPEK SVWTAPAQPSGPSPELQGQRSR RNGWSWPPHPLQIVAWLLYLF FAVIGFGILVPLLPHHWVPAGY AVSLGKRRAGPWNPFLSVSAET ILPQCMGAIFAGHLVVHLTAVS IDPADANVRDKSYAGPLPIFNR SQHAHVIEDLHCNLCNVDVSA RSKHCSACNKCVCGFDHHCKWLNNCVGERNYRLFLHSVASALL GVLLLV/LGGHICLRGVLCQPH ASAHQPTL
22892	53260	A	23024	2	224	OODECEC AL ADCECT DDDDCDI
22893	53261	A	23025	[11]	1368	QQPSGSGALARGSGLPPRPGRL PARSPVQKRKGGWPKGKKRNP PRDLAV\RVPTTGYMIFLNEQRS QLRATHPDLPFTEIMKMLAVQ WAQLSQDKKGRKYAGKALGV TPPPSALTPVALIEEPKQLDIQSL PFQPILDRLKIKDKRQLVPSYLG GMHTVTEGLAGRQVLLQQVQ GGAWESAFLTCPWAMAAAAP GPTLTMLQPSISAVGFSQMAMG PHVQGRASKLSIYVNPRRQLST TTAMGDELNDLYCGTSRQLFSS LCNKKEPAAKAAPAAPLSSPGP PALGHAGHGRQLLLERSLTLFH LYELLKYPRASPGLWPPRVWM PHSLTCGTFWVPMIMMRFCAH VASGLLRPPFNQVFAHLSPAQG RHPGENGIIVLTSRAAVRIRCSS CSRSLTGNLKKILPRTQSTWSL WRKRRDK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22894	53262	Α	23026	2	405	
22895	53263	Α	23027	1	391	
22896	53264	Α	23028	3	463	
22897	53265	Α	23029	74	466	FAFNMPEPAKSAPAPKKGSKK AVTKAQKKDGKKRKRSRKESY SVYVYKVLKQVHPDTGISSKA MGIMNSFVNDIFERIAGEASRL AHYNKRSTITSREIQTAVRLLLP G\EL\AKHAVSEGTK\AVTKYHQ
22898	53266	A	23030	168	597	KEETDFLFWFCHYCFIMPELAK SAPAPKKGSKKAVTKAQKKDG KKRKRSRKESYSVYVYKVLKQ VH\PDTGISSKAMGIMNSFVTTF FE\RIAGEASRLAHYNKRSTITS REIQTAVRLLLPGELAKHA\VSE GTNA\VTKYTSSK
22899	53267	A	23031	18	321	FTATWLNQEVGIVGKYRTRYG VSLQKMVKKIEISQYAKYICSFL WQNSKMKRRAVGIWCCG/SCM KTVKVSSSRNHIVWTYNTTSAV TVKSTIRRLKQLKDQ
22900	53268	A	23032	12	445	FTCRSHVSEVEERCVEAPGLLG SRSITGFSLHSKQAKRTKKVGIV \GKYGNPLWGPSLR\KMVKK\IE ISQHAQVHFALFCGKTKMKRR AVGIWHCGSCMKTV\AGGAWT \YNTT\SAVTVKSAIRRLKEFERP SRTLLYSLRTSLGL
22901	53269	Α	23033	3	175	
22902	53270	С	23034	378	434	
22903	53271	A	23035		578	MVRCVRLVEAGSVVRYLSTSIC RPVVDAGSRALCLQEWADSQQ VKEKQYSSRDVQRAAALNIYRI PPSSRKPALCPTPRDRLEYDEDR LEHIAYVRARELHTLEVTGLET VAQSKAHVASLEGLIPEDKVVL LAGSPLQNEATLGQCGVEALTT LEVVGRRLGASL/HTSASKHTM VRALTYCSSREETFTAI
22904	53272	C	23037	236	379	
22905	53273	A	23038	178	534	VVQKTEPRLCFAVSPPFSCSLPL PSLFPHSPLLGRPKRQAKPAAD EGFWDCSVCTFRNSAEAFKCSI CDVRKGTSTRYLKICVTWVVV VI*QVFIEHTAICALGVQWYKD RPGPCSYGA

SEQ ID	-	l .	SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,21/	sequence	or peptide sequence	defection,possible nucleotide insertion)
22906	53274	A	23039	44	1012	 VGGGGVQPSMTMGDKKSPTRP
						KRQAKPAADEGFWDCSVCTFR
						NSAEAFKCSICD\VRKGTSTRKP
						RINSQLVAQQVAQQYATPPPPK
						KEKKEKVEKAGQKRKPEKRPR
•						KFSP/SVVTKKNTNKKTKPKSD\
						ILKDPPS\EANSIQSANATTKTSE
						TNHT\SRPRLEKRGTGGIAQQFG
						QLTVG\NVTVIITDFK\EKTRSSS\
						TSSSTVTS\SAGS\EQQNQRQLR
						GFRGATDKGSLPVPPNPKGRPC
						QAVNDESFLKLAHGNCGKLWN
1						SGVWKFQNLPPAHACLHPWEN
					1	LSCGTSTSLVDCCQDNFLLCHG
						HSWPPRDFRHPDDYSGHF
22907	53275	A	23040	1	626	MAGCRNPCKAESHRSKSLLSEA
ŀ						ESLFREAESQHDRGSKNAGHR
		ĺ				DESVEVWLSLLSSSQTRAEIER
						QSRGKQWLRHLLRDTEGPKET
						VRPELATRAAMDVAEVEFLAE
						KELVTIIPNFSLDKSYLIGVDNIP
						KADEIRTLVKDIWDTRIAKLRV
	1					SADSFVRQQEAHAKPDNLTLM
						EINPSGTFLTQALNHTYK/LRTN
						LQPSESTQSQDF
22908	53276	Α	23041	1	951	MDAAEVEFLAEKELVTIIPNFSL
						DKIYLIGTWELKYREIRWLSQG
						PHGKPHADPGSSILQNSSGPSG
į						VRTNGKSKPWEPFVAEEFAHQ
						FHESVLQSTQKALQKHKGSVA
						VLSAEQNHKVDTSVHYNIPELQ
						SSSRAPPPQHNGQQEPPTARKG
						PPTQELDRDSEQEQEEDDEDGE
						DEEEVPKRKWQGIEAVFEAYQ
						EHIEERTLERNRSSTAADKAQL
						LAANRQGHGQPTDPASDNIPKA
-						DEIRTLVKDMWDTRIAKLRVS
ĺ						ADSF\VRQQEAHAKL\DNLT\LM
						EIHTSG\TFLTQALNHMYKLRT
22000	52077		22042	1	(2)	NLQPLESTQSQDF MAGCRNPCKAESHRSKSLLSEA
22909	53277	Α	23042	1	626	
						ESLFREAESQHDRGSKNAGHR
Ì						DESVEVWLSLLSSSQTRAEIER
						QSRGKQWLRHLLRDTEGPKET VRPELATRAAMDVAEVEFLAE
						VRPELATRAAMDVAEVEFLAE KELVTIIPNFSLDKSYLIGVDNIP
						KADEIRTLVKDIWDTRIAKLRV
						SADSFVRQQEAHAKPDNLTLM
]						EINPSGTFLTQALNHTYK/LRTN
	l		L		L	LQPSESTQSQDF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22910	53278	A	23043	2	642	RPRRLLRDAEGPEETVRLWPAA RAAMDAAEVEFLAEKELVTIIP NFSLDKIYLIGGDLGPFNPGLPV EVPLWLAINLKQRQKCRLLPPE WMDVEKLEKMRDHE\RKEETF TPMPKPFTTLELT\KLLLNHASD NIPKQDEIRTLVKDMWDTRIAK LRVSADSF\VRQQEAHAKL\DN LT\LMEIHTSG\TFLTQALNHMY KLRTNLQPLESTQSQDF
22911	53279	A	23044	3	1038	TSSMAAGCSEAPRPAAG\SDGS LVGQAGVLPCLELPTYAAACA LVNSRYSCLVAGPHQRHIALSP RYLNRKRTGIREQLDAELLRYS ESLLGVPIAYDNIKVVGELGDIY DDQGHIHLNIEADFVIFCPEPGQ KLMGIVNKVSSSHIGCLVHGCF NASIPKPE\QLSAEQWQTMEIN MGDELEFEVFRLDSDAAGVFCI RGKLNITSLQFKRSEVSEEVTEN GTEEAAKKPKKKKKKDPETY EVDSGTTKLADDADDTPMEES ALQNTNNANGIWEEEPKKKK KKKH\QEVSRTQDPCFSQGQLT S\SGYQSDHKKKKKKKRKHSEEA
22912	53280	В	23045	82	323	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
]	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
22212	150001	<u> </u>	100046		1.604	OVER A DEVINOR A DAY A
22913	53281	Α	23046	3	1624	QLEIRAPTADEIHVTVGEARNLI
		ŀ				PMDPNGLSDPYVKLKLIPDPRN
		1				LTKQKTRTVKATLNPVWNETF
						VFNLKPGDVERRLSVEVWDWD
						RTSRNDFMGAMSFGVSELLKA
		ŀ				PVDGWYKLLNQEEGEYYNVPV
		ļ				ADADNCSLLQKFEACNYPLEL
						YERVRMGPSSSPIPSPSPSPTDPK
						RCFFGASPGRLHISDFSFLMVLG
}		1				KGSFGKVMLAERRGSDELYAI
						KILKKDVIVQDDDVDCTLVEKR
		l				VLALGGRGPGGRPHFLTQLHST
		ŀ				FQTPDRLYFVMEYVTGGDLMY
				1		HIQQLGKFKEPHAAFYAAEIAI
						GLFFLHNQGIIYRDLKLDNVML
						DAEGHIKITDFGMCKENVFPGT
						TTRTFCGTPDYIAPEIIAYQPYG
			,			KSVDWWSFGVLLYEMLAGQPP
		l				FDGEDEEELFQAIMEQTVTYPK
		l				SLSREAVAICKGFLTKHPGEAP
						GASGP*WGNLTIRAHGFFPLGF
		ŀ				DWERLERLEIPASFSRPRPCGRS
		ĺ				GENFDKFFTRAAPALTPPDRLV
						LASIDQADFQGFTYVNPDFVHP
						DARSPTSPVPVPVM
22914	53282	Λ	23047	1	1522	MLRAPRQATRDHHFAHIDIVQV
		l				TCFRYLRRSALAPAPAMPSPVR
		1				VDRGSFPPVAGGNHTRRRGGTP
						QEGVTREALAFFDVVATTGAE
		l				MMWSNFFLQEENRRRGAAGRR
		l				RAHGQGRSGLTPEREGKVKLA
						LLLAAWAPRWRCCPWAPSSG
						WTQHLQGQRQRRVRSGPTWG
		1				CGRRAPSGCGRRTCPWTGTPA
		l				APRSCPEASERNSEHPALICTLP
		1				QPHMLKRRARRAPGSMAPPLP
		l				RPPQAASVPVFPPDWEPLESRA
	1	l				RSKSPLHSQHRPAQETGRHSRP
						RTSTRKRQQKHSTEKATHTPER
						APKPQTRENNPKGRAQGSPAPR
						DHNPASRHQKTSNDKRSTHDT
						HKTRKDSQTRRKKTRNNEKQQ
						PDR/IGPQIIAYQP/YGKSVDWW
						SFGVLLYE/MLAGQPP/FDGEDE
						EELFQAIMEQ/TVTYPKSLSREA
						VAIC/KGCGRSGENFDK/FFTRA
						APALTPPDRLV/LASIDQADFQG
						FTYVNP/DFVHPDARSPTSPVPV
						PASSPTRPQLLENKFGC
L	<u> </u>	J	<u>L</u>	<u> </u>	L	LYPPLIKI GEFFINKLAC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22915	53283	A	23048	505	2345	SIWGMVMMNPTPSSITLRRMM
22713	33203	``	25040		23.13	SLFPASLTTKYGGFYINSGTLQF
	ļ					RQASESEDDFIKEKKKKSPKKR
						KLKEGGEKIKKKKKDDTYDKE
						•
						KKSKKSKFSKAGFTALNASKEK
						KKKKYSGALSVKEMLKKFQKE
						KEAQKKREEEHKPVAVPSAEA
						QGLRELEGASDPLLSLFGSTSD
						NDLLQAATAMDSLTDLDLEHL
						LSESPEGSPFRDMDDGSDSLGV
						GLDQEFRQPSSLPEGLPAPLEKR
						VKELAQAARAAEGESRQKFFT
						QDINGILLDIEAQTRELSSQVRS
		1				GVYAYLASFLPCSKDALLKRA
						RKLHLYEQGGRLKEPLQKLKE
		1				AIGRAMPEQMAKYQDECQAHT
		1				QAKVAKMLEEEKDKEQRDRIC
		1				SDEEEDEEKGGRRIMGPRKKFQ
						WNDEIRELLCQVVKIKLESQDL\
		1				ERNNKAQ\AWEDCVKGFLDAE
		1				VKPLWPRGWMQARTLFKESRR
		1				GHGHLTSILAKKKVMAPSKIKV
		1				KESSTKPDKKVSVPSGQIGGPIA
		1				LPSDHQTGGLSIGASSRELPSQA
ŀ						SGGLANPPPVNLEDSLDEDLIR
		l				NPASSVEAVSKELAALNSRAAG
						NSEFTLPAPSKAPAEKVGGVLC
						TEEKRNFAKPSPSAPPPASSLQS
22916	53284	Α	23049	1	379	AYRNFTSASDVWSYGIVMWEV
						MSYGERPYWDMSNQD\PNTAL .
						LDPASPEFSAVVSVGDWLQAIK
						MDRYKDNFTAAGYTTLEAVVH
						VNQEDLARIGITAITHQNKILSS
						VQAMRTQMQQMHGRMVPV
22917	53285	Α	23050	1	123	
22918	53286	Α	23051	3	221	
22919	53287	C	23052	80	379	
22920	53288	A	23053	71	924	AFSVNSSNSWEKA*SQKCLQV
						NYLSLSSASYLP/CMTCPPSAPL
						NLISNVNETSVNLEWSSPQNTG
						GRQDISYNVVCKKCGAGDPSK
						CRPCGSGVHYTPQQNGLKTTK
						VSITDLLAHTNYTFEIWAVNGV
						SKYNPNPDQSVSVTVTTNQAAP
						SSIALVQAKEVTRYSVALAWLE
						PDRPNGVILEYEVKYYEKDQNE
						RSYRIVRTAARNTDIKGLNPLTS
						YVFHVRARTAAGYGDFSEPLE
						VTTNTGNKDHPDSVTSEAIFIIQ
			1		-	LNPITVKSKELSVIFGSADILLQ
L	<u> </u>	ل	I	I	<u> </u>	DITTIT TRURDED TH GUADIELQ

SEQ ID	SEQ ID NO:		SEQ ID NO:	1	L .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,21/	sequence	or pepade sequence	deterion, (-possible nucleotide ruser tion)
				,		
22921	53289	Α	23054	1	1102	MRWLLKWERKYTLYSSGVLK
						HKNLLESLLGQFLGPPAQRFDS
						AGGVRPKNDGRFTVIQLVGML
						RGIGSGMKYLSDMSYVHRDLA
	-					ARNILVNSNLVCKVSDFGMSR
						VLEDDPEAAYTTRGGKIPIRWT
						APEAIAYRKFTSA\SDVWSYGIV
						MWE\VMSYGERPYWDMSNQD
1						VIKAIEEGYRLPPPMDCPIALHQ
1						LMLDCWQKERSDRPKFGQIVN
1						MLDKLIRNPNSLKRTGTESSRP
1						NTALLDPSSPEFSAVVSVGDWL
						QAIKMDRYKDNFTAAGYTTLE
l						AVVHVNQ/EVSTQRCNTKGDL
ľ						ARIGITAITHQNKILS\SVQAMR
İ						TQMQQ\MHGRMVPRLSTVLNK
	1					LKTLEISLPHPCTLIEELHFFNFV
22922	53290	Α	23055	148	3229	NAFRFHLLNPAGESHDRLQDLL
ł						SAVRSPTKRQEQRWHRRTMAG
						IFYFALFSCLFGICDAVTGSRVY
						PANEVTLLDSRSVQGELGWIAS
						PLEGGWEEVSIMDEKNTPIRTY
		1				QVCNVMEPSQNNWLRTDWITR
						EGAQRVYIEIKFTLRDCNSLPG
		ŀ				VMGTCKETFNLYYYESDNDKE
						RFIRENQFVKIDTIAADESFTQV
1						DIGDRIMKLNTEIRDVGPLSKK
		1				GFYLAFQDVGACIALVSVRVFY
						KKCPLTVRNLAQFP
22923	53291	Α	23056	57	450	VGDCLTSRGMSWVQATLLAR/
						ALCRAWGGT/CRGALTGTSISQ
						VPRLAP\RGLHCSA/ASVSSEQS
						LVPSPPEPRQR/PPGGERDKASF
						LQTVQKFADTSVVDSGHIDFIY
					1	LALRKMREYGVERDLAVYNQL
		<u> </u>				LNIFP

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22924	53292	ĪA	23057	89	1502	VGDCLTSRGMSWVQATLLARG
						LCRAWGGTCGAAL/SQEPPSLR
Ì						SLASLPRGLHCSAAAHSSEQV\L
		1				VPSPPEPRQEGPPRLLVPFEGPV
						LGQGAGLGERGQGRAFL/LDGC
						RNLRTTSVRKRGPH/VDLFYLA
					•	LRKMRE\YGVERG/DLAVYNQL
						LNIFPKEVFRPRNIIQRIFVHYPR
						QQECGIAVLEQMENHGVMPNK
		ŀ				ETEFLLIQIFGRKSYPMLKLVRL
						KLWFPRFMNVNPFPVPRDLPQD
						PVELAMFGLRHMEPDLSARVTI
	i					YQVPLPKDSTGASKIPPKPHIVG
					ļ.	HRACMSPHQESKKSRSSKAAL\
						ARHNPS\RPVF\VEGPFSLWLRN
						K\CVYY\HILRA\DLLPPEEEGK
						WKRTPEEWE\LYYP\RQLDLEY
						VRSG\WDNYEFDINEVEEGPVF
						AMCMAGAHDQ\ATM\VKWI\Q
				-		GPAGDQPQTLAQNPPWSFRLRR
						VQPGELPDILLQGWRSP/HPAR
		1		}		GPPRKKTTTCKRQQQGPELV
22925	53293	Α	23058	1	2517	
22926	53294	Α	23059	1	927	IIVVEFVSRGLARAPSGPAVFSH
						LAMARLPLRGVRWGCLLTAVH
Ì						PEPPTACREKQYLINSQCCSLCQ
						PGQKLVSDCTEFTETECLPCGE
						SEFLDTWNRETHCHQHKYCDP
						NLGLRVQQKGTSETDTICTCEE
						GWHCTSEACESCVLHRSCSPGF
						GVKQIATGVSDTICEPCPVGFFS
						NVSSAFEKCHPWTSCETKDL\V
					ŀ	VQQAGTNKTDVVCGPQDRLRA
	1					L\VVIPIIFGILFAILL\VLVFIKKV
						AKKPTNKAPHPK\Q\EPQEINFP
						DDLPG/SSNTAAPVAGRIFTWG
		<u> </u>		<u> </u>	100	GQPV\TQEDGQRESHFSCRERQ
22927	53295	В	23060	24	439	
22928	53296	A C	23061	1	582 2607	
22929	53297		23062 23063	248	430	ATNYTREQDSMWNCVR*GRTT
22930	53298	Α	23003	240	730	ATTHSVTQRTDERQRRADTAL
						TLADYRNTSVTGVFAGLF
22931	53299	A	23064	776	1426	I LEAD I KIN 13 V I U VI TAULI
22932	53300	A	23065	284	598	VFRLLRGQKSEPGKSRPKAL*N
22932	33300	^	23003	204		KPANTPVQSHISLYAAGIINSDS
				1		KVR*SCRQSSARQSGSYL*TSPR
						SPQNRQQTGEITATVIAFGRTRR
						TAPDITPHPASAR
22933	53301	A	23066	3139	3792	III DIII III ROFII
22933	199901	T.,	23000	3133	3132	<u> </u>

SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>		sequence		
22934	53302	Ā	23067	2	830	GVPNMAAPQDVHVRICNQEIV
						KFDLEVKALIQDIRDCSGPLSAL
					:	TELNTKVKEKFQQLRHRIQPVL
			:			YQRAFIWTASTFFFKLTYSLTDF
		ł				SSTQHDFNSPTTPVTFSDL\EQL
		l				AKEQDKESENQLLLQEVENHK
						KQMLSNQASWRKANLTCKIAI
						DNLEKAELLQGGDLLRQRKTT
						KESLAQTSSTITESLMGISRMM
		1				AQQVQQSEEAMQSLVTSSRTIL
		l				DANEEFKSMSGTIQLGRKLITK
						YNR\RELTDKLLIFLALRLFLAT
2222		<u> </u>	220.60		1450	VLYIVKKRLFPFL
22935 22936	53303 53304	A B	23068 23069	1231	1458 2133	
22937	53305	A	23070	722	957	VFRLLRGQKSEPGQKQAKSTLE
22931	33303	^	23070	1/22	931	QTSEYPSNRGIPHRQSECSIRTPL
						PFVRALRD*VRGCGCAPLSSVD
		1				SFFCCTEAL
22938	53306	A	23071	2	1137	VPNMAAPQDVHVRICNQEIVKF
22/30	33300	ľ``	25071	[11137	DL\EVKAPIQDIRD\CSGPLS\AL
						TELNPKVKEKFQQLRHRPTGP/
		l				VEQLAKEQDKESEKQLLLQ\EV
						ENHKK\QMLQQFRPSWEGKLN
						SSPGKICNSDNLEKAELLQGGD
						LLRQRKTTKE\SLAQTSSTIT\ES
						LMG/ISGRMMAQQVQQSEEAM
						QSLVTSSRTILDANEEFKSMSGT
						\IQLGRKLITKYNRRELTDKLLIF
						LALALFLATV\LYIVKKRLFPIF
					ŀ	VRFPKVPVLAPFNSLFSGICPWF
						PEAPRLPKAEAHTPLRFAPQLP
						CRLGWEHQCPPLFLQSSWLVK
						GEQSPWEEGPGIGMPPLGTYEP
						PPSHHCALWEAAAAWSTLKSE
						PPRTRMPLLSPAPIVPLNARSGG
						GKREER
22939	53307	Α	23072	64	665	RVRGGEAELAMAAAEEE\SGAP
						NVPNRERGGAGRPKTPFECNIC
						LETAREAVVSVCGHLYCWPCL
						HQWLETR\PERQECP\VCKAGIS
						REKVVP\LYGRGSQEAPRIPELK
	1					TPPPPPRGQR\PAPESRGGF\QPF
						GDTGGFHF\SFGVGAFPF\GFFT
						TVF\NAHEPFRRGTGVDLG\QG
						HPSLQLGKEFPLPGFSAIFFLFW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22940	53308	Ā	23073	1	416	MQRRRRRRLLLLPPRRQTDRAL
22710	33300	1	25075	<i>'</i>		PPPPSPGSMPPPLPPPP/PPARGG
		İ				GLRPGDGLYPK/PSPARRGSGA
						RTSWCTMCAPPSTSAPRASRRP
						RPSSCATRPPTSKPPPA\GHAPH
						PPPRDLGGGLDSGVQSDGVLQ
						HLQRPGH
22941	53309	Α	23074	1	1713	TIEQUE ON
22942	53310	A	23075	1305	1538	LKVKRNCTSS*SPTIGHD*QSNL
227.2						SLPPSDFAKPPFPFSLLRCSKFR
						KLKAAYRKRKKGHKSSLSLSV
						KINKTAAANK
22943	53311	Α	23076	49	355	
	53312	В	23077	253	471	
22945	53313	Α	23078	1267	1497	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
						KLKAAYRKRKKGHKSSLSLSV
						KINKTAAANK
22946	53314	Α	23079	3	1237	HEVAGAVPGAIMDEDYYGSAA
						EWGDEADGGQQEDDSGEGED
						DAEVQQECLHKFSTRDYIMEPS
						IFNTLKRYFQAGGSPENVIQLLS
						ENYTAVAQ\AVNLLAE\WLIQT
		ŀ				GVEPVQVQDTVEYHLKSLLIKH
						FDPRIADSMFTEEGETPAWLE\Q
1						MIAHTTWRDLFYKLAEAHPDC
						LMLNFTVKLISDAGYQGEITSV
						STACQQLEVFSRVLRTSLATILD
						GGEENLEKNLPEFAKMVCHGE
						HTYLFAQAMMSVLAQEEQGGS
						AVRRIDQEVQRFAQEKTVLTQE
	:					AIITVKGVSLSSYLEGLMASTIS
						SNASKTIVAAMSNLVPPVELAN
			}			PENQFRVDYILSVMNVPDFDFP
						PEFYEHAKALWEDEGVRACYE
						RSNEYQLIDCAQYIFVLEMQVP
						CCEEAQSHPGQHGGVQVGRN
22947	53315	Α	23080	1358	1589	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
		l				KLKAAYRKRKKGHKSSLSLSV
						KINKTAAANK
22948	53316	Α	23081	33	458	GITGSTDASGVTMKFNPFVFSD
						RRKNRKRHVNAPSHVRRKIMS
						SPLSKELRQKYNVRSMPIRKDD
						EVQVVRGHYKGQQIGKVVHV
						YRKKYVIYIERVQREKANG/DK
						DRKKILERKAKSRQVGKEKGK
						YKEELIGKMQEYIEP
22949	53317	Α	23082	1	576	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22951	53319	Α	23084	124	549	NNSLSSPVQADVYFPRLIVPFCG
						HIKGGMRPGKKVGVMGIVDLN
						PKSFAINLTCGDSEDPPADGAIE
						LKAVFTDRQLLRNSCISGERGE
•		İ				EQVSNPLLSIHSTTMPFRVEILC
		İ				EH\HCFR\VFVDGHQLFDFYHPH
						SNRYLAI
22952	53320	Α	23085	1	215	QGFIKPPLKEENFTLPPPGKFGP
						PKGSLNSPPS/PSSP*SPGGGILLL
						VYYLYRALFQKGCKEAAWLEY
		l				PLT
22953	53321	Α	23086	533	983	PWGPGCNKRWPCLAKDPPPTP
						QSSGVGQPFGGAEPAGAYIRTG
						PHLGPPPPPPPDSSTNLHSPNPS
						DDGADTPLAQSDEEERGDGG
						AEP\G\PAASSGPLAYRPDPAGY
						SPTSRPPGPSPEPLG/AEDPDSLL
	ł		İ			ASRAPRGGNGWWDCGTLS
22954	53322	Α	23087	1422	1670	NRYIDKTGWAWWLTPVIRHFR
						RPRWVDCLRLGVREQPGQHSKI
						KSLQKNAK\ISQAWWHAPVDP
	İ	l				ATWKADAGGSLEPGSLRLQ
22955	53323	Α	23088	2	663	VLLDERSAALDGAKRDGTLAL
		1				AAGALCREARAAQVFFLKGGY
						EAFSASCPELCSKQ/INVSANCP
						NHFEGHYQYKSILCGMTTHKA
	1		!			DISSWFNEAIDFIDSIKNAGGRV
1						FVHCQAGISRSATICLAYLMRT
ţ						NRVKLDEAFEFVKQRRSIISPNF
		1				SFMGQLLQLESQVLAPHCSAEA
İ						GSPAMAVLDRGTSTTTVFNFPV
				L		SIPDHSTNSALSYLQSLITTSSHC
22956	53324	Α	23089	3	206	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22057	52225	 	23090	982	2252	LFLEKRRPALGPENRQRAPQGR
22957	53325	Α	23090	982	22.32	STLGALRPAMVMEVGTL\DAG
						1
						GLRAL\LGERAAQLPACLDCRS FF\AFNAGHIAGSVNVRFSTIVR\
]					
		l				RRAKG\AIGALE\HIVPNAELR\G
						RLLAGAYHAVVLLDERSAG\LD
						GAKRDGTLALAAGRA/LCREA
1						RAAQVFF/LSKGGYEA\FSASCP\
		1				ELCSKQSTPMGLS\LSLSTSVPD
						SAESGC\SSCSTP\LYD\QGGPVE
		1				ILPFLYLGSAYHASRKD\MLDA
						LGHRLPWINVLSQIVPNHF\EGH
						\YQYKSIPVEDNHKADISFWFN\
						EAIDFIDSIKNAGGRVFVHCQA
						GISRSAT\ICLAYLMRTNRVKLD
						EA\FEFVKQRRSIISP\NFQLHGA
						SLLQFEFPRMLGPALFRAEAGK
						PRQMAVL\DRGT\STTTVFNFPV
						SIPVHSTNSALSVPFRAPLTDLS
						QLVERPTGR
22958	53326	Α	23091	13	287	
22959	53327	Α	23092	1	492	
22960	53328	A	23093	1	636	
22961	53329	Α	23094	1	506	PTRTWTRGRIPRLSAQPSRGAR
						GTMADPRVRQIKIKTGV\VKRL
						V\KEKVLYEKEPKQ\QEEKIEKM
				-		RAEDGENYDIKKQAEIL\QES\R
						MMIP\D\CQRR\LEAAYLDLQRI
						LENEKDL/EKEAEEYKEAR\LVL
				1		DSVKLESLKLFSYGVVFALKPR
						GPILQSIIFDHVLCVQV
22962	53330	Α	23095	1	786	
22963	53331	Α	23096	183	1059	VLGAFIWQGLDPSPETAFTDHR
]				1		ATEWGTNDQSLCLNSP*\DAPS
						AKHEYQASVSGSVSSKPEIDLQ
						LYDEVRDQDVPVKRNFARIVV
						NVSDTNDHAPWFTASSYKGRV
		1				YESAAVGSVVLQVTALDKDKG
	·					KNAEVLYSIESGSHPPYKTAAD
						VLLKAPPPGWRPTNTKPVHSTE
		1				AQPTQNQRTQQKHNQGASRSP
						RHSFATSTRAGAGICGCKTRRW
		1				MTRWMEESRGSLQTLPSTRLEP
		1				GSSAAWLDPEEQAQSLQFSSQE
1						APFLGEGGENHIKGVPHGTKES
[QQQPLNHRSSL
22964	53332	Α	23097	1	1224	
22965	53333	Α	23098	1	660	

SEQ ID	SEQ ID NO:	1	SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	l	09/340,217	sequence	or peptide sequence	deterior, —possible nucleotide insertion)
		l				
22966	53334	Α	23099	129	595	LFPRLLSCLTTPPHCSFSICFVIC
						SRTLILKGSSLMYVFCLPNTAIV
						MALSPRGWRSKFGMPWHSKPT
	İ	ł				WIMTL*FPLEKTGALSLTSDTS
		l				MFTVVVLTLGGLPLSEA*T/GG
						NYDSHFDVDKGTGTIIVAKPLD
1	İ					AEQKSNYNLTVEATDGTTTILT
22967	53335	Α	23100	1	1782	MAARIEVIHELSIRLLLMKTDL
	ĺ					DAATAALLNVQPAKSGDNTSG
						APFPGKISQTLLKDPHNGYHVV
	ŀ					FALAGSPKDADDTSIYMFYQK
	Ì					VGDNSIDSWKNAGRVFKDSDK
						FDANDPILKDQTQEWSGSATFT
		}				SDGKIRLFYTDYSGKHYGKQSL
	1					TTAQVNVSKSDDTLKINGVED
						HKTIFDGDGKTYQNVQQFIDEG
		ŀ				NYTSGDNHTLRDPHYVEDKGH
i						KYLVFEANTGTENGYQGEESLF
						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTFEKSNEAADHFKHGGGATIK
						GRVADNGKPQLSSLTYIDIRVIE
			1			ESIYPPAILPLEIFITASGEEYSGG
		ŀ	:			VIGKIHATDQDVYDTLTYSLDP
						QMDNLFSVSSTGGKLIAHKKLD
						IGQYLLNVSVTDGKFTTVADIT
	:					VHIRQVTQEMLNHTIAIRFANL
			į			TPEEFVGDYWRNFQRALR\NIL
						GVRRNDIQIVSLQSSEPHPHLDV
		l	•			LLFVEKPGSAQISTKQLLHKINS
		l				SVTDIEEIIGVRILNVFQKLCAG
						LDCPWKFCDEKVSVDESVMST
						HSTARLSFVTPRHHRAARQQ\R
						GPPLHNPADT*TTARKYNCDKR
22968	53336	Α	23101	184	13958	AMGRHLALLLLLLLLFQHFGDS
]				DGSQRLEQTPLQFTHLEYNVTV
}						QENSAAKTYVGHPVKMGVYIT
1						HPAWEVRYKIVSGDSENLFKAE
						EYILGDFCFLRIRTKGGNTAILN
						REVKDHYTLIVKALEKNTNVE
]						ARTKVRVQVLDTNDLRPLFSPT
]		1				SYSVSLPENTAIRTSIARVSATD
]		1		1		ADIGTNGEFYYSFKDRTDMFAI
						HPTSGVIVLTGRLDYLETKLYE
						MEILAADRGMKLYGSSGISSMA
						KLTVHIEQANECA
22969	53337	С	23102	56	331	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22970	53338	A	23103	2249	2680	GEFLKTREFSIYFLEKYTTFPGE NLEHGFWDSSTSPPVLLHPGIPA SRI/PVSSIFTKVPPKKSATKP/PV SIWGAPPLGPPGFWGCCPTGEQ RGVCHPSEPHPPVVDFRKKPPF LCVSLSLINIFKSNKFLLTTCNL VIFYTCSL
22971	53339	A	23104	739	1672	CQEAASEFGGPLHTPAMFLRRL GGWLPRPWGRRKPMRPDPPYP EPRRVDSSSENSGSDWDSAPET MEDVGHPKTKDSGALRVSRAA SEPSKEEPQVEQLGSKRMDSLK WDQPISSTQESGRLEAGGASPK LRWDHVDSGGTRRPGVSPEGG L\GVPGPGAPLEKPGRREKLLG WLRGEPGAPSRYLGGPEECLQI STNLTLHLLELLASALLALCSRP LRAALDTLGLRGPLGLWLHGL LSFLAALHGLHAVLSLLTAHPL HFACLFGLLQALVLAVSLREPN GDEAATDWESEGLEREGEEQR GDPGKGL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22972	53340	A	23105	122	2437	RLELKLPLWTMRFTEMELLGH LSIFKITGKQKQLTRTAPGPSGQ LRSNQTTGNSLEPSGQPRSSQTT GNSLEPSGQPRSSQTTGNSLEPS GQPRSSQTTGNSLEPSGQPRSSQ TTGNSLEPSGQPRSSQTTGNSLE PSGQPRSSQTTGDSLEPSGQLRS NQTTGDSVQPNNREQPGTVRS APLQPNNREQPGTVRSAPLQPN NRGQRPTKQQGTTWNRQVSSA PTKQQGTAR\PSGQPRSNQTTG NSLEPSSQLRSNQTTGDSVQPN NREQPGTVRSAPLQPNNRGQRP
						TKQQGTAR\PSQPCSNQTTGN SLEPSSQLRSNQTTGDSVQPNN REQPGTIRSAPLQPNNRGQRWD RQVSSAPTKQGTALGPSG\SSAP TKQQGTALEPSSQLRSNQTTGN SLEPSGQPRSNQTTGNSREPSSQ LCSNQTIGNSQDRQVSSAP/IQT TGNSQDRQVSPAPTKQQGTAG NRQVSST/PIQTGNSVQPNNREQ PGTVSGAM/VWVATPVKGPAR VETSAHKVSAAGLPKKKT/QPG VTGGAPRLTLNTVLPKHSRTPS RHEAGSTQALYHNGQAQYALS FPHEAAPMAAFYVVTGIEVTTA LTLEEAPCTVEITTALTLEEPPC TVFWNKTERNTRTLVQDKGSN TKGFVQTATCCFRADCKASRPS
22973	53341	A	23106	605	813	PHPPSPHREWTSDLRHKQVSKF PGAHRLPGHPELPIQEAWDRGE LFVFINFRENPALLRW\AYAR\TI NVYP\NFRPTPKNSLMGALCGF
22974	53342	A	23108	273	936	GPLIFIYYIIKTERVSIQTRCLVF AYSNKVPSWNQKA\PSPTKPAG TLIFDFPAVKTVRNKFVFFINYP VSGYTEDLKKFLKPYTLEEQKN LTVCPDGALFEQKGPVYVACQ SPISLLQACRGMNDLDFGYSQR NPCILVKMNRIIGLKSEGVPRID CVSKNEDIPNVAVYPHNGIIDL KYFPYYGEKLHVGYLQPLVAV QVSFAPNDTGKEVTVECKIDGS ANLKSQDDRDKFLGRVMFKIT
22975	53343	A	23109	104	478	ETEVGGLVEASMSHNCTTALQ PGQHSQKENLQNKPIPGREKY* VSIEFELIYFP/YYCQVGYLQPL VAVQVSFAPNNTGKEVTVECKI \DGSANLKSQ\DDRDKFLG\RV MFQNHSTVHSMSRVSSTE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22976	53344	Α	23110	3	903	LLSPVRAHTMTKNEKKSLNQSL AE\WKLFIYNPTTGEFLGRTAAS WGLILLFYLVFYGFL\AALFSFT MWVMLQTLNDEVPKYRDQIPS
						PGLMVFPKP\VTALEYTFSRSDP TSYAGYIEDLKKFLKPYT\LEEQ KNLTV\CPDG\ALFEQKGPVYV ACQFP\ISLLQACSGMNDPDFG
						YSQGNPCILVKLNRIIG\LKPEG VPRIDLCFKRMKDIPNVAVYS\ HNGMIDLKYFPYYGKKTAMF/
						VGIYQPIGLLVQVSFAP*QPLGK EVTVE/CAKIDGSSQPKKVQDD\ RDKVFGDEFMFQIHSTVH
22977	53345	A	23111	467	931	SRDSRRGTPLAGGRGRDFAGQ MGHRVQGRHGSRGRLPSGDAR VNAERAEPVGAGAGGPGGSS VRGFCGRSWSEA/RGRRRAGG
						GVH\IEP\RYRQFP\QLTQIPRCSQ SEF\FSGLMWFWIF\WRFWHDP K\KVLGHFPV/YPDPSQWTD\EE L\GIPPDD
22978	53346	A	23112	1	390	VTIPTSLAELQGLPHIRVFLDNN PWVCDCHMADMVTWLKETEV VQGKDRLTCAYPEKMRNRVLL ELNSADLDCDPILPPSLQTSYVF LGIG*ALIGAIFLHYRNNGPLTIS SEGGLASLLCKDSRGVPLL

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22979	53347	A	23113	3	1653	PQRIQRGVPTRGEEVAPGRRQE EEREQERGAERPDPPCVLSGGK
						GRGNRPLREAPGGEGVSQVPA
						AAPLPRFHERKVFFFQTLPPAR
						ALRAQPPEPSERAPSQPSSGETR
						AAMPGGCSRGPAAGDGRLRLA
						RLALVLLGWVSSSSPTSSASSFS
						SSAPFLASAVSAQPPLPDQCPAL
						/CACRSEAARTVKCV\NRNLTE
						VPTDLPAYVRNLFLTGNQLAVL
						PAGAFARRPPLAELAALNLSGS
						RLDEVRAGAFEHLPSLRQLDLS
						HNPLADLSPFAFSGSNASVSAPS
						PLVELILNHIVPPEDERQNRSFE
						GMVVAALLAGRALQGLRRLEL
						ASNHFLYLPRDVLAQLPSLRHL
			į			DLSNNSLVSLTYVSFRNLTHLE
						SLHLEDNALKVLHNGTLAELQ
						GLPHIRVFLDNNPWVCDCHMA
						DMVTWLKETEVVQGKDRLTC
						AYPEKMRNRVLLELNSADLDC
			,			DPILPPSLQTSYVFLGIVLALIGA
		1				IFLLVLYLNRKGIKKWMHNIRD
						ACRDHMEGYHYRYEINADPRL
		ļ .				TNLSSNSGCLEEILEDRPRTTLH
22980	53348	Α	23114	288	877	ASRLSLPTRTSMRSSSSTSPSSSS
			!			SSSSLCRFLLNSRVTDAAFNFLL
						VWYYCTLTIRESILINNGSRWG
						RAGPEGGEHGMGCQGPSGKGR
						EQGAGG*WWVFCSRV\KGWW
						VFHHYVSTFLSGVMLTWPDGL
						MYQKFRNQFLSFSMYQSFVQF
						LQYYYQSGCLYRLRALGERHT
						MDLTVGESGVALASGHSPFLPS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22981	53349	Α	23115	Ti	1273	MAPLHSILGNRVGLCQASVRA
						CPGATVSSQCRASLRSRLAWRG
						SELPANRGRTWPGPGGLRLQA
						APRLAYLGRATLLVLTPYYDLP
						PNVRFTSMPASPHTYFLVLGAPI
						FLLLRSTWSYYHACRFEYATPN
						DRNTTYLAISGPTRRTYDVATF
						AYKDEYEKFKLYLTIILILISFTC
						RFLLNSRVTDAAFNFLLVWYY
						CTLTIRESILINNGSRPDGLMYQ
		İ				KFRNQFLSFSMYQSFVQFL\QY
						YYQSGCL\YRLRALG\ERHTM\D
						LTVEGFQSW\MWRGLTFLLPFL
		ŀ				FLW/YNSGQLFNALTLFKPGPR
						DPQLQGVGRCLCCGFPFLL\LFL
		ľ				GNFFTTLRVVHHKFHSQRHGS
	•	İ				KKDLRLGLSPCRPRRGFLVPVC
			İ			FWEGDWEGAPSSVRVSGGLFY
						SPLGFMGRCGPGKEDPGPSCPQ
22982	53350	A	23116	238	398	
22983	53351	Α	23117	1	386	
22984	53352	A	23118	13	580	
22985	53353	В	23119	39	1171	
22986	53354	Α	23120	381	795	PPSSKAPSSKGPAKIVCIYWALI
				1		LQSNEVTITEDKFNNLIKAAAV
		ł		1		TVEPFWPSFFAKALASVNIGSLL
						CNAGVG/SMAPAACTAQAGGP
]		ATSTTAASASFRNHNPRIAPKFR
	ŀ					GVLAHEQYLSERATWVWAPNP
		İ		4		NPKSLF
22987	53355	Α	23121	1	960	
22988	53356	Α	23122	2	3952	
22989	53357	Α	23123	1058	1494	KSVRLVVN/YLRTQKAVVRVSP
	1	ĺ				EVPLQNILPVICAKCEVSPEHVV
						LLRDNIAGEELELSKSLNELGIK
		l				ELYAWDNRRVPSAVPMQGPWP
						AEHFLQVNEQPPAGSPSPCPSSS
		ŀ				SSSSGLRGGMQTANGPACALTP
		ŀ				TCAQTDCPCPTSH
22990	53358	Α	23124	195	398	TDSSSALQPQHRGRPALRQRTP
		l ⁻				PRRSP/SPGLSPPRGASPAS\SGA
		l				PARGVAFGGDEPPIGAEAMYSK
						P
22991	53359	Α	23125	2	155	*
22992	53360	A	23126	1	339	
22772	199900	Ľ `	123120	I *	555	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22993	53361	Α	23127	150	1017	LGRWRQDQVPHGLECRSQSEW
						SCTDSRQHSSS\YSDSRRQEIGK
						MRYVSVRDFKGKVLIDIREYW
		İ				MDLKRLPRKPRTGEAGSGSLSS
1						PGADTPVEAAVTWRLGLGSVD
		ł				VGSFRAWQRGKSAAVGHTVSP
						RWPRWQGGDSSALSMASLSTL
						TCSWGLRYSALSPTTSAQREEG
}						SALYRHLSPPSLRFYPTVKWAT
					ļ	LAITRPRPPAPRSHSHHPQPRAL
						CPVWVLFQHPKTHGENLGDRW
						KEGSALVSSKGTPAKLSSGRIR
						NALQNTLGEVSVARIQTHRGIF
						VETTEARL
22994	53362	A	23128	l .	584	APIECGGIPSLPVLCQSRANDQE
		l				GVRLLEPSEAMPKSKELVSSSSS
l				}		GSDSDSEVDKKLKRKKASLL/P
						ENPVKK\QKTGETS\RALSSF*TE
						PASSRDDNMFQIGKM\RYVSV\
		l				RDFKGKSA*FDIREYWMDPEGE
İ		l				MKPERKGISLNP\QQWEPSLKE
		l				QISGPLIDASKKTCKISEPILIKPC
						TVPVVLICLVTVAVCF
22995	53363	Α	23129	186	687	LLWARGLGRAKSAVPTVSTML
						GLPWKGGLSWALLLLLLGSQIL
		l				LIYAWHFHEQRDCDEHNVMAR
						YLPATVEFAVHTFNQQSKDYY
						AYRLGHIFNSWKEQVESKTV\F
ļ						SMELLLGRTRVWEYLKTTIDNC
		l				HFQESTELNNTFTCFFTISTRPW
						MTQFSLLNKTCLEGFH
22996	53364	Α	23130	232	402	LFKAWECFFLIFLQVVITRLKLD
						KDRKKILERKAKSRQVGKEKG
						KYKEETI\EKMQE
22997	53365	A	23131	637	758	NVEEKNRKKILERKAKSCQI\K
						KEKGEYKEETIGTMRECGL

Sequence Sequence	,
22998 53366 A 23132 1 1095 IVFLLYLETC LSDSSERLFSI DGFNFQGTRI GAIEDWNNE IDGYNAQYN FKRRKGPVHI REYLTHSKLM ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPF ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	FGVIADVQFADLE RRYYRHSLLHLQ SSMPCCVLQLGDI ASKKSLELVLW\Q HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
LSDSSERLFSI DGFNFQGTRI GAIEDWNNE IDGYNAQYN FKRRKGPVHI REYLTHSKLN ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPF ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	FGVIADVQFADLE RRYYRHSLLHLQ SSMPCCVLQLGDI ASKKSLELVLW\Q HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
DGFNFQGTRI GAIEDWNNE IIDGYNAQYN FKRRKGPVHI REYLTHSKLN ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPI ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	RRYYRHSLLHLQ SSMPCCVLQLGDI ASKKSLELVLW\Q HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI ALGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
GAIEDWNNE IDGYNAQYN FKRRKGPVHI REYLTHSKLM ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPH ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 15552	SSMPCCVLQLGDI ASKKSLELVLW\Q HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
IDGYNAQYN FKRKGPVH REYLTHSKLM ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPH ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE CONTROL	ASKKSLELVLW\Q HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
FKRRKGPVH REYLTHSKLM ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPI ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	HTWGNHEFYNFS NTKFLEDQIVHHP AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
REYLTHSKLM ETMPSEDYY LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPF ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	NTKFLEDQIVHHP AYHFVPFPKFWFI LLGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET
ETMPSEDYY, LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPH ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	AYHFVPFPKFWFI LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
LLDAYDLS/A GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPH ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE	LGAWDQSSPKYG SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
GSVWKILEGS SRDFSEA/RSI LNWLNEVLT HLP\LYPGPPI ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	SHNP\NTGNWNSS LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
SRDFSEA/RSI	LFQFNGRVSAQEQ FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
LNWLNEVLT HLP\LYPGPPH ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	FSDTNQEKVVIVS FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
HLP\LYPGPPH ALAVIWSHEG GGYSEDPFG APDSQRFGT RGRVPDRIYE	FDNVCLAWNYRD CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
ALAVIWSHEG GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	CV\GVSLLGHTHD VYHG\NLEGVIET VHVYPDKMMLKG
GGYSEDPFGV APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	VYHG\NLEGVIET VHVYPDKMMLKG
APDSQRFGTV RGRVPDRIYE 22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	VHVYPDKMMLKG
RGRVPDRIYE	
22999 53367 C 23133 235 382 23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	ELQGKEEPSHC
23000 53368 B 23134 63 293 23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	
23001 53369 A 23135 160 330 23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	
23002 53370 B 23136 1 870 23003 53371 A 23137 2 1013 23004 53372 A 23138 1 1552	
23003 53371 A 23137 2 1013 23004 53372 A 23138 I 1552	
23004 53372 A 23138 1 1552	
23005 53373 B 23139 120 1034	CLLLACDCAAAA
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	FLLLAERGAAAA RARLGCLPGGLR
1 1 1 1 1	LTSCIDPSMGLNE
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	FDFAAREMAPNM
	PVDVMRKAAQL
1 1 1 1 1	DVGGSGHVRLDT
1 1 1 1 1 1	GCTSTTAY\IRHPQ
1 1 1 1 1 1	DEGGTHYKAGHL
	CLPPGRPCHRGA
23007 53375 A 23141 2 412	- CELT GIG CINCON
23008 53376 A 23142 47 487	
<u> </u>	GRCLEAQGVINY
1 [] 1 [] 1 [] 1	VLVAHPVF*SFSCE
1 1 1 1 1	MTMLHCRHSDPV
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	DFASVVVLCPPPQ
	FGGLPLSIDFVMT
VDFCCHLHLI	1
	AGQQVLDVNSCL
1 1 1 1 1	TFSGLHAE/AEGS
	LGLKTLCTSQQAF
	LDDSKSDLEEPW
ELRKEGAPRS	1
23011 53379 A 23145 1 207	
23012 53380 A 23146 l 3009	
23013 53381 A 23147 3 298	
23014 53382 A 23148 1 636	
23015 53383 A 23149 1 795	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23016	53384	Α	23150	125	1527	SSPRQQTQLARVHP/DGRAQ/LA
						PSCLPGLGLCPLPVCPPVCNCPA
				1		CPCHEHCGEAP*GVA\AKRGAL
		1				RVHPHPSCGGPRAMTVTLHGQ
					Ì	DVLMLPGDLVGLQHDAGPGAL
					ļ	LHCSPAPGHPGPQAPYLSANAS
		ŀ				SWLPHLPAQLKGTWACPACAL
						RLLAATEQLTVLLGLRPNLGLR
	ļ					LPGRYEVRAEVANGVSRHNLS
						CSFDVVSPVAGLRVIYPAPRDG
						RLYVPTNGSALVLQVDSGANA
		Ì				TATARWPGGSVSARFENVCPA
						LVATFVPGCPWETNDTLFSVVA
						LPWLSEGEHVVDVVVENSASR
						ANLSLRVTAEEPICGLRATPSPE
						ARVLQGVLVVLLAGSSGYLVG
						FKFLESHGSDSGSANSFHCLISQ
İ						NEFKTLLPDLTRAPRYSPVVEA
						GSDMVFRWTINDKQSLTFQNV
•					-	VFNVIYQSAAVFKLSLSADSAP
						RPRWRLQSPRVGRLSPDPGFFG
						SPGPCQPRLQAFLI
23017	53385	A	23151	648	1387	ASTQSLLAWCPAWLSIPSTWPS
					}	SFSSGCPG/CKAQPVYEVSRSSY
						VYLEGRCLNCSSGSKRGGYTFT
						LT\VLGRSGEEEGCASIPLSPNRP
						PLGGSCRLFPLGAVHALTTKVH
						FECMGWHDAEDAGAPLVYAL
						LLQRCRQGHCEEFCVYKGSLSG
						YGAVLPPGFRPQFEVGLAVVV
						QDQLGAAVVALNRSLAITLPEP
						NGSAMGLTVWLHGLTASVLPG
1						LLRQADPQLVIEYSLALVTVLN
						ETYTESL
23018	53386	A	23152	2	654	DKHIWLSIWDRPPRSCFTRIQRA
						TCCVLLICLFLGANAVWYGAV
						GDSAYSTGHVSRLSPLSVDTV\
						AVGLVSSVVVYPVYLAILFLFW
						MSRSKV\AGSPSPTPAGQQVLDI
	1					DSCLDSSVLDSSFLTFSGLHAEV
						INTLADHQHRGTDFGGSPSVLII
	1			-		TVSLRSYKFAISLCTTYLWVINT
						LADHRHRGTDFGGSPWLLIITV
						FLRSYKFAISLCTTYLC
L	L	<u> </u>	l	<u> </u>	L	I DAOTAI AIGEOTTIEC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23019	53387	Α	23153	2	4072	SRDPAGSYHLNLSSHFRWSALQ
						VSVGLYTSLCQYFSEEDMVWR
				1		TEGLLPLEETSPRQAVCLTRHL
						TAFGASLFVPPSHVRFVFPEPTA
						DVNYIVMLTCAVCLVTYMVM
						AAILHKLDQLDASRGRAIPFCG
						QRGRFKYEILVKTGWGRGSGT
						TAHVGIMLYGVDSRSGHRHLD
						GDRAFHRNSLDIFRIATPH\SLG
						SVWKIRV\WHDNKGLSLALGFL
						Q\HVIVRDLQTA\QSTFFLVNDW
			ļ ļ			LSVETEANGGLVEKE
23020	53388	Α	23154	1	10922	MPPAAPARLALALGLGLWLGA
						LAGGPGRGCGPCEPPCLCGPAP
						GAACRVNCSGRGLRTLGPALRI
						PADATALTAMAKADGDVKKY
İ						GSKRGALALTHGHSLLRDVSH
ŀ						NLLRALDVGLLANLSALAELDI
						SNNKISTLEEGIFANLFNLSEINL
						SGNPFECDCGLAWLPRWAEEQ
	ł					QVRVVQPEAATCAGPGSLAGQ
						PLLGIPLLDSGCGATLVGPHGPL
						ASGQLAAFHIAAPLPVTATRWD
						FGDGSAEVDAAGPAAS
23021	53389	Α	23155	3887	14184	HVLCAWVTYMVMAAILHKLD
						QLDASRGCAIPFCGQRGRFKYE
						ILVKTGWGRGSGTTAHVGIHA
						VWGGQPERPPAPGTATEPSTAT
						VWDIFQIRHPAQPGVACWKIRV
						FGTTTKGSALPRFPCSTSSSGTC
		i				RRHAAPSSWSMTGFRWRRRPT
						GAWWRRRCWPRVRPRSVFPLR
						HAALLRFRRLLGAELQRGFFDK
						HIWLSIWDRPPRSYFTHIQRATC
						CVLLICLFLGANAVWYGAVGD
		_				SAYSTGRVSRLNPLSVY
23022	53390	Α	23156	1	442	GGASGMGGIT\ASTVNQ\SLLSP
						LVLEVDP\NIQAVRTQEKEQIK\
						TLNNKFASFIDKVRFLEQQNKM
						LETKWSLLQQQKTARSNMDN
						MFESYINNLRRQLETLGQEKLK
						LEAEFTTWGPPLPMPPATKQFN
		l				CFFFLVQNKTSASSANVK
23023	53391	A	23157	1	497	
23024	53392	С	23158	208	422	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23025	53393	A	23159		1339	ELSSPRSSFSAPSRISAWFGPPAS TPASTMSIRGTQKSYKGSTSGP RAFKQAAS*QRSGPGFRISSSSF SRVGSSNFRGGLG\GGYGGASG \MGGITAVTVNQ\SRALLSPLVL EVDPNIQAVRTQEKEQIKTLNN KFAS\FI\DEVGFLEQQNKMLGT KWSLLDQQKTA\RSNM\DNMF\ ESYINNLRRQLETLAQEKLKLE AELGNMQGLVEDFKNNYED\EI NKSTETENEFVLIKKDVDEAYI NELYEEEIRELQSQISDTSVVLSI DNSRSLDTDSIIAEVKTEISEMN RNISWLQAETEGLKGGRISLEA GIADAEQRRELVIKDANAKLSE LEAALQQTKKLLE\MENMSIHT KTTSGYAGGVSLAYGGLTGSG LSYGLSSSSRSGAGSSSFSRTSST RAMVVKKIKTRDGKLLTLSLR
23026	53394	A	23160		651	MTAFNSGKVDIVAINDPFIDLN YMVYMFLYDSTHGKFHGTVK AENGTLVINGNPIT\IFQDQDPS KIKWAP/LAKVIHDNFGIIEGFM TTVHTITATQTINGPSGKLSRDG CRALQDIIPVSTGAAKPVGKVIP ELNGKLTGMAFHVPTANVSVA DLTCRLEKPA/KYDDIKK\NTHS STFDAGAAIVLKDHSVKLISWY DNEFGYSNRVVHLMAHNASKE
23027	53395	В	23161	1	382	
23028	53396	A	23162	3	1151	ASAHASAVRQSAAIFFCVASRA HIAETPLGKVKVGVNGFGRIGR LVTRAAFNSGKVDIVAINDPFID LNYMVYMFQYDSTHGKFHGT\ VKAENGKLVINGNPITI\FQE\RD PSKIKWG\DAGAEYRRGSPLGV FNQPWRKAGGSICKGGSQKGSS ISAPSAECPPCFVMGCEPMKKY DKQAFKI\ISN\ASCTNQLL*HPL AKG*FHDNFGNSLEGLHDHKS MAITANPRKTVD\GPSRGKLWP LMGRGAL\QNIIPASTG\AAKAV GKVIPELNGKLTGMAFRVPTAN VSVVDLTCRLEKPAKYDDIKK VVKQASEGPLKGILGYTEHQV VSSDFNSDTHSSTFDAGA\GIAL\ NDHFVKLISWY\DNEFGYSNRV V\DLM\AHMGFQGS
23029	53397	A	23163	160	236	
23030	53398	Α	23164	931	1124	
23031	53399	Α	23165	132	303	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23032	53400	IA	23166	20	171	
23033	53401	A	23167	1008	1262	HELQCSLLPVRHATAGHTNGQ
		Ì				NRLHWGHVAHSNYGAALKTN
						GVEVR*FTAAIARCDGNPYESF
		l				RKDVRADFQTRIDAIVRCLPK
23034	53402	A	23168	732	1162	RCHEKGRKGHQAPTGSTRPDG
2505.	33 102	-	23.00	1.52		TLSACRSCQQGQGMPASAGVP
		1				ARQMGSTGNLPHLV*PYFLGLS
						SKAQHTG*LHKKR
23035	53403	A	23169	2	1093	
23036	53404	Α	23170	236	496	
23037	53405	В	23171	22	3084	
23038	53406	A	23172	305	522	LRDGRGSMSSSSPNSASGSAICP
						ASSVMPETVSVSRVMPNVTKP
]	ŀ					A/MTAKQTRIRNQMVAITYIES
1	į					DAWNRSYQ
23039	53407	A	23173	177	386	
23040	53408	A	23174	3	200	GAVAPMHPLAHPVPGAAAQPP
	Į.					PA*QPATPCHQPA*GQPAPPDV
		ĺ				PGEPSAGRSLHHAHICLAPNEQ
23041	53409	Α	23175	171	535	
23042	53410	A	23176	190	505	VPTDRIRTRLTSVCKRRCEVHQ
						LEDRPFEVYQVA/PRRD*SFETS
		1				VWNHMRKQ/PPLDVRQLAS*N
						VLHRQLDIMDSAVASMYSIHLS
						IKMGIGFSGVTTVVKGTPK
23043	53411	Α	23177	349	1272	
23044	53412	Α	23178	518	1451	VSSSIIMFGPLSQFPRQQVGSSV
						KRPSNVVCPGLMFSFFSSAA*IP
						**PLI*QPGV
23045	53413	Α	23179	2436	2894	TSPDVLRAGPRVPSVQGTSPSQ
						SCPGQRPHGPNSKAYGR/LSVM
1		1				AQYYCNVIPVLE/VPPSAFTPPP
		1				KVDSAVV/RLVPHATMPHPVK
		1				DVRVLSRITTEAFNQRRKTIRNS
	1					LGNLFSVEVLTGMGIDPAMRA
	1					ENISVAQYCQMANYLAENAPL
23046	53414	A	23180	1846	2557	IPVSERQGSDWRFFSVYVVNKA
						GGLIYQLDSYAPTGLRLEKTF\S
	İ	ŀ				YPLDLLLKLHDERVLVAFGQR
						GRHPEWGHAVL\AINGMDVNG
						RYTAD\GKKVLEYLG\NP\ANYP
						V\SYLFAQPRL\TSNDKLMLAL
						MFHSLFAIGSQACLP\EQGKPQG
						IENAWKTDTFKLH\CYQTLT\GI
						KFVVLADPRQAGIDSLLRKIYEI
						YSDFALKNPFYSLEMPIRCELFD
						QNLKLALEVAEKAGTFGPGS
23047	53415	c	23181	34	225	
23048	53416	В	23182	105	381	
23049	53417	A	23183	1	837	
<u> </u>		1.,	1	<u> </u>		<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
-			:	sequence		
23050	53418	Α	23184	3	744	HEASCRSERRRQMADFDTYDD
1						RAYSSFGGGRGSRGS\AGGHGS
1						RSQKELPTEPPYTAYVGNLPF\N
		l				TVQGDIDAIFKDLSIRSVR\LVR
				•		DKVTGKFK\GFCYVEFDEV\DSL
						K\EALTYD\GALLGDRSL\RVDF
						AKGRKTKIRVAFGF\RKGGPDD
						R\GFRDDF\LGGRGGSRPGD\RR
						TGPPMGSR\FRD\GPPLRG\SN\M
						DFREPTEEERAQRPRIQ\LKPRT
					•	VWRPLNQVANPNSAIFGGARPR
						EEVVQKEQE
22051	52410		22105	1	1038	EEVVQREQE
23051	53419 53420	C A	23185 23186	1	1486	MGRNQSRKAENSKHESTYSPP
23032	33420	A	23180	l l	1400	·
				İ		KDHSSSQAMEQSWTENGFEKL
						GFRRKEALYYLDLTASCQSQEL
	ŀ	İ				FQLYAGMSVVGTSMPVQAVCP
						YCGNRIITVTTFVPGALTWLLC
			1			TTLFLFGYVLGCCFLAFCIRSLM
	1					DVKHSCPVCQRELFYYHRLRQ
	ļ	ŀ				MELASSARTVVAPSRPCAGREG
						RGDANGKNSSPTAGSAMSSEPP
			<u> </u>			PPPQPPTHQA\GVGLLDTPRSRE
						RSPSP\LRGNVVP\SPLPTRRTRT
1						FS\ATVR\ASQGPVYKGV\CKCF
			ŀ			CRSKG/HIGFNYPPAGWAGPDIF
						PGTSLDVGRGSY\VPSGKGDEV
						TLLKCAPIPTPKNE\KLAGPWEV
			}			VHHSPWHPGT\KHET\WSG\HVI
						SFLGGLVESTPLSLCFVGDFAG
						RRQQTLEDDILPHETGLQRGNG
			ŀ			PSQVSPGGKGYGGAGVGCGVF
						PAISTAYGPLQQPLHHLKSIKSH
						LKRRGAHWWLSGGSNPIPGSGS
						RVGISSQLAQTHGLNPQNPSSS
						WSWRPQIPRSGPLAA
23053	53421	Α	23187	2	617	
23054	53422	A	23188	1	993	
23055	53423	Α	23189	779	1210	AAAGSEAGRRRGADPTRGPHV
						LPGSRQRRP/PIFPSLDLSGFFLF
						CSEFCPEIKSTNPGISIGDVAKE
						KKKLGEMWNSLNDSEKQPYFT
]			KAAKLKEKYEKDVADYKSKG
						KFDGAKGPAKVARKKVEEEDE
						QEEEEEEEEEEDE
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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23056	53424	A	23190	116	522	KAPYSTNSLSLPPFPILFMGSGY RKGPTFQEPKPLPGAAARGPGC DPTQPPRHSAPRDHGPP\SPRCS PRPARPRASPAWLPPRAPTPPSP
						PAPAAAPRSCRGHARASSGRSG TTATPRRRRGGLWNQSPMDIK G
23057	53425	Α	23191	1	152	
23058	53426	A	23192	1	676	MWSAGRGGAAWPVLLGLLLA LLVPGGGAAKTGAELVTCGSV LKLLNTHHRVRLHS\HDI\TYGS GSGQQ\SVTGVEASDDANSYW RIRGGSEGGCPRGSPVRCGQAV RLTHVLTGKNLHTHHFPSPLSN NQEV\SAFGE\EGEGDDPGTLW TVRCSGTA/HWEREAAVRFQH VGTF\VFLSVTGEQYGSPIRGQH EVHGMPQCQ\QHNTWKAMEGI FIKPSVEPSAGHDEL
23059	53427	A	23193	1	1044	
23060	53428	A	23194		1083	MTATLAAAADIATMVSGSSGL AAARLLSR\SFLLPHNGIRHCAY TA/SLRQHLYVDKNTKIICQGFT GKQG/TPFHSQQALEYGTKLVG GTTPGKGGQTHLGLPVFNTVK EAKEQTGATG\SVIYVPPPFAAA AINEAIE\AEIPLVVC\YPRGIPQP GLGHSSSHKTCGPGKDKAYWG PNCPGV\INPGECKIGIMPG/HIF HKKGRIGIVSTSGTLTYEAGH/Q TTQKLGLGQSLCVGIGGDP\FN GTDFID\CLEILFERFLPQEGIILI G\EIGGNA\EENAAEILEPLYSGS KF/ARPVVS\FIAGLTAPPGRRN GVMPGAIISGREKVGAKRERSS ALQSAGVVV\SMSPCTAGEPRF YKEFEKREDRL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
23061	53429	Α	23195	1173	2110	FFTELEETILKFIWNQKRACISK
						ARVNTNDKSGGITLPDFKLYNE
						AIVAKTTWYWYKSRHIDQEKR
						REEKKE*KKDPSVPETLKKKQR
						NFTELKIQCLSKKFAQKMLQTA
						RRKLIYEKVKHCHKEYRKMYR
						TEIRMARMATKAGNFYVPAEP
<u> </u>						KLAFVIRIRGINGVSPKVRKVLQ
						LFLLYQIFNGAFVKLNKASINM
						LRIVEPYIA/WRYPNLKSVNELI
						YKCGYGKINKK*IALTDNALTA
						RSLGKYGIICREDLIHEIYTVGK
		1				CFKEANNFLWPF/K/LSSPRGGM
						KKKTTHFVEGGDAGKREVQIN
						•
23062	53430	A	23196	3	571	RLIRRMN GTMEGVEEKKKEVPAVPETLK
23062	33430	A	23190	3	3/1	l I
ŀ					1	KKRRNFAELKIKRLRKKFAQK
						MLRKARRKLIYEKAKHYHKEY
					[RQMYRT\EIRMA\RMARK\AGN
						F\YVPARTQIWRFVIR\IRGINGS
						EPQRFRKVLQLLRPFVKIFNG/T
				1		PFVKLQQGFRFNMLRDCRSHIL
						AWG\YPN\LKSVNELIYK\RGYG
22052	50.401	ļ	22107		2.40	QNQFDLHFPTEKRSHPGSQ
23063	53431	Α	23197	2	340	PRVRPRVRDSKMASVGEVSGP
				1		QYPVKDKKLL\EVQTGGSFPSW
l						\ILMRGPSSPSGIFG\AFQRGLLR
						GTYNKYI\NVKKGSDLGGFTMV
		ł				L\ACY\VLFSYSFSYKHLKH\ER\
		ļ.,	22100		10	LRKYH
23064	53432	Α	23198	2	48	PRVRLRC*MVDEYCGCEWGK*
		ļ.,_				LNGGRVLWM
23065	53433	A	23199	123	693	QACIMREYKLVVLGSGGVGKS
						ALTVQFVQGIFVEKYDPTIEDS
						YRKQVEVDAQQCMLEILDT\AG
		İ				TEQFTAMRDLYMKNGQGFALV
ļ	ł					YSITAQSTFN\DLQDLREQILRV
						KDTDDVPMILVGNKCDLEDER
]						VVGKEQGQNLARQWNNCAFL
						ESSAKSKIYVNEIFYDLVRQINR
						KTP\VPG\KAPQKSPSCQLL
23066	53434	Α	23200	461	1422	SPHGEPLLGQCRREMWGWSLH
						TESPLG\QWGSQILKLQNDLL
23067	53435	Α	23201	232	331	TPL*KWLRGVMAVTVVTPQTG
						IIAIPDPHHEWGSTGYPRLPA

SEQ ID	1 7		SEQ ID NO:	E .	I .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23068	53436	Α	23202	5	1223	GLDSGPESGLRAQPGLLAGKRK
						MSVLRRMMRVSNRSLLAFIFFF
1						SLSSSCLYFIYVAPGIANTYLFM
		1				VQARGIMLRENVKTIGHMIRLY
						TNKNSTLNGTDYPEGNNSSDYL
						VQTTTYLPENFTYSPYLPCPEKL
						PYMRGFLNVNVSEVSFDEIHQL
						FSKDLDIEPGGHWRPKDCKPR
						WKVAVLIPFRNRHEHLPIFFLHL
						IPMLQKQRLEFAFYVVEQTGTQ
						PFNRAMLFNVGFKEAMKDSV
						WDCVIFHDVDHLPENDRNYYG
						CGEMPRHFAAKLDKYMYILPY
						KEFFGGVSGLTVEQFRKINGFP
						NAFWG\WGGEDDDLW\NRV\H
						YAGY\NV\TRPEGDLGKYKSIPH
						HHRGEVQFLGRYKLLRYSKER
1						QYIDGLNNLIYRPKILVDRLYT
ĺ						NISVNLMPELAPIEDY
23069	53437	Α	23203	78	495	CTFEKAPPFQLGRGRMVPAKK
						GGEKKTGR\AINEVVTREYTINI
İ		Ì				HKRIHGVGFKKRAPWALKEIR
ĺ						KFAMKEMGTPDVRIDTRLNKA
ĺ			ĺ			VWAKGIRNVLYRIHVRLSRKR
ĺ						NEDEDSPNKLYTLVTYVTVTTF
ĺ	ļ					KNLQTVNVDEN
23070	53438	Α	23204	138	360	KNQKRQLSGGHCADQTSLVLA
[Ì.	RLAPLF\PTTMSDKPDMAEIEKF
1						DKSKLKKTETQEKNPLPSKETT
	l					EQEKQAGKS
23071	53439	Α	23205	145	728	KRQPTSAMKDPSRSSTSPSIINE
						DVIINGHSHEDDNPFA\EYMWD
						GKMEGRNFNR\QIGRRELWGR
					<u> </u>	KD\FLNRCFPKE\MLGEEGKEG
						ALNWVLFPSFEILPHQTMGPNP
		ŀ	<u> </u>			RTSFNDPCLSSDGSFSWEDLVV
	1					KS\NLNPNAKEFVPGGEVRKYL
						SRRGPLLVDVAQFPHCEGSIRR
l				İ		LNCKSSLGHCVTLYAIAQSFC
23072	53440	Α	23206	530	707	
23073	53441	A	23207	1	515	MDFSKVPPHAAFRAQPPPALT
		1				WGLAKGSGRGAGEGGERPDRR
						R/PAARTGTPAWRPAGRRRLPP
						RSPPPAPVPVRPALPHPVRPGKE
						TGKTRRRHYSPGRPLRGAAA
		1				AAAAAAVAQTASPSWAEGGRS
		•				GAGGSAESPAFSSPALPCPEGLL
		1				CPSAVPVPSKQSVSEVLAIF
23074	53442	Α	23208	3	713	
23017	22112	11.	122200	12	1/13	

SEQ ID	SEQ ID NO:	i .			ř .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23076	53444	Α	23210	2	1631	GLPRRPTRPPTRSGVSPPWPGPH
						SGVLFPLLLHCTGSLSQAVLTQ
			;			PSSLSASPG\ASVKLTCTLRSDIN
						VGSPFIYWYQQKPGSPPQFLLR
						YKSDSDNQQGSGVPSRFSGSKD
		•				ASANAGILLISGLQSE\VRDDYY
						CQAWDSSTAHSDTGRCGIVGRP
						PVRQCWPLLLLGLAMVAHGLL
						RPMVAPQSGDPDPGASVGSSRS
						SLRSLWGRLLLQPSPQRADPRC
				}		WPRGFWSEPQSLCYVFGTGTK
			}			VTVLGQPKANPTVTLFPPSSEEL
						QANKATLVCLISDFYPGAVTVA
						WKADGSPVKAGVETTKPSKQS
ļ						NNKYAASSYLSLTPEQWKSHR
						SYSCQLQWGLGLGASQGGFLY
						EPVSQCVVFGGGTKLTVLGQP
						KAAPSVTLFPPSSEELQANKAT
						LVCLISDFYPGAVTVAWKADSS
						PVKAGVETTTPSKQSNNKYAA
•						SSYLSLTPEQWKSHRSYSCQKP
						NHASHGTSRMPSVSAPVSGLKS
						CWSGTGDRSALGQMPGPCHGI
						PDPMQQASDSGQPKAAPLGSL
						CSKPSSEELSGQQGPHWCVS
23077	53445	Α	23211	1	959	VFLLLYLVAMPDPAKSAPAPK
						KGSKKAVTKAQKKDGKKRKR
						SRKESYSVYVYKVLKQVHPDT
						GISS\KAMGIMNSFVNDIFERI\A
						GEASRLAHYNKRSTITSREIQTA
						VRLLLPGELAKHAVFIENLLNY
ļ						RKYLVKSSEKDRNYPCPLPGLK
						SKVDICTPYCLLLLPQPRHYPG
						KDTKEETEDFPNSPPPINREEDK
						RYTTVMGPCLRQAALEGELLA
						CPVMEDQQANWVHEPITFNTF
						KEIRKSIRENGPASPFTRGLIEVI
						ADSSDLVASTQKLTQPKETPSA
	1					PYNDISDLTNQDSWFIGFPPIWN
22070	50446	ļ. —	02010		163	AGETDLSNNKTT
23078 23079	53446	A A	23212 23213	3 74	463 464	
23079	53447	A	23213	41	490	
23080	53449	A	23214	2	369	PDPAANAPAAAKPAAAKALPV
23001	33447	(``	-3213	[~		FLLLLSGGAGPAHPMRAPGC*G
						ELGYTEQLFQFLQATAQEEGIF
						EYTDPKLAISAVWTFRDLGVLQ
						QTPSPAGPRLHLSPTFASLDNQE
						KLEQFIRQFICS
23082	53450	A	23216	124	301	112241 11441 100
25002	UUTUU	14	23210	147	201	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23083	53451	Α	23217	3	2426	PGVLSTATFRESGLSPCSKLKHP
						GPRGRLQYSPPTAGQGRLENG
						AGHPQSTGPVATLSVAEYGTQ
						GGASGPARIPIGSRGCLLAGRA
						EASLWSSGFGMKLEAVTPFLG
						KYRPFVGRCCQTCTPKSWESLF
						HRSITDLGFCNVILVKEENTRLF
						SWALLRVLKLLFLNVQLHKGQ
						MKMVQKAAQQVGQVGTSLWL
						SPEQGLPLVLLSTHKTLLDGILL
						PFMLLSQGLGVLRVAWDSRAC
1						SPALRALLRKLGGLFLPPEASLS
						LDSSEGLLARAVVQAVIEQLLV
1	ľ					SGQPLLIFLEEPPGALGPRLSAL
1						GQAWVGFVVQAVQVGIVPDAL
1						LVPVAVTYDLVPDAPCDIDHAS
						APLGLWTGALAVLRSLWSRWG
						CSHRICSRVHLAQPFSLQEYIVS
						ARSCWGGRQTLEQLLQPIVLGQ
						CTAVPDTEKEQEWTPITGPLLA
ŀ						LKEEDQLLVRRLSCHVLSASVG
	}					SSAVMSTAIMATLLLFKHQKG
		ŀ				VFLSQLLGEFSWLTEEILVRGFD
		ŀ				VGFSGQLRSLLQHSLSLLRAHV
						ALLRIRQGDLLVVPQPGPGLTH
						LAQLSAELLPVFLSEAVGACAV
						RGLLAGRVPPQGPWELQGILLL
					:	SQNELYRQILLLMHLLPQDLLL
1			 		:	LKPCQSSYCYCQEVLDRLIQCG
						LLVAEETPGSRPACDTGRQRLS
						RKLLWKPSGDFTDSDSDDFGE
						ADGRYFRLSQQSHCPDFFLFLC
23084	53452	A	23218	169	955	ASSQLPCSMAQCCATWAPLKM
						RWLCSTAWSTSSLSLPPSWSSL
						QASQPSCCHATSLAPPCAGQCL
						ARAWPVLSFLCTCALGLLASIA
						MTFATQGKALLAACTFGSSELL
				:		ALAPYCPFDPTRIYSSSLCLWGI
						ALRCSAWAENVFAVRCAQLTH
						QLLELRP\WWGKKQPTHMMRE
						NPELVEG\RDLLSCT\SSE\PLTL
						CRDDVLPRP\DGQLRTLQATLL
						CDPGPGFLGAGLRGLKWTLGG
						P\SGAFQPSSPTTPAHLHLKIETL
						F
23085	53453	Α	23219	1	344	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23086	53454	Α	23220	214	1355	FCSSWLRSLADSSLSWKMFLVG
		-				LTGGIASGKSSVIQVFQQLGCA
						VIDVDVTARHVVQPGYPAHRRI
						VEVFGTEVLLENGDINRKVLGD
						LIFNQPDRRES\LNAITHPEIRKE
						MMKETFKYFLRGYRYVILDIPL
						LFETKKLLKYMKHTVVVYCDR
1		1				DTQLARLMRRNSLNRKDAEAR
		l				INAQLSLTDKARMARHVLDNS
			İ			GEWSVTKRQVILLHTELERSLE
						YLPLRFGVLTGLAAIASLLYLLT
				ļ		HYLLPYAYTASAIPPSGAASRG
				1		PGARCSAIRKRAPARPTSAVSPP
ŀ						GAPRSPARGLRAIDRAAGTRNL
						NATLPPPGPPAPPSPPSPPSSITL
						HVFHQLQPDITRGREGTDFQWE
						KFPGRVLVADSRRALRCLAPAP
ļ		l				TWLV
23087	53455	Α	23221	2	1231	TPLFLRKGTADGSARTTVIHLG
		}	<u> </u>			FTKGKQRCVETQAAAVAGDVG
			1			RTAAGGRVGKESGVISSDDRPG
			1			LPVGQSQPEGGEQEATDTVSVH
						ATTSQAWPGGADSKELSSKAE
			1			ERETGKDQQTELPLCLPLRDFP
			İ			DSAPVMSSDSGHATESLHLASS
						DSKPPMPTPNEGPGSTVGHAGP
						PRTVKADLCTEPQLALRGLQVS
						LPRLHLVGVIYAEHQASGLEGC
						GHAGSCSPDGSPRAPGATLCLS
						KPLSFRTKRCLPCWLPASAPVGI
						PGAPVHIPGAPVCIPGAPACISN
						APVHIPGAPVGIPGAPVCIPGAA
						VCVPNTPMCIPSAPMCIPNTPVC
						IPNTPVCIPNTPVCIPGAAVCILG
			1			A/PRAHPWCSSCVHPGAPECYL
						GCGLTQADRLEPAVLDAGLQE
		<u> </u>				PLFALCGQALPS
23088	53456	A	23222	2	408	
23089	53457	Α	23223	62	251	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1
ł	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23090	53458	Α	23224	2111	3003	TQKAICMCLGLCKKIHHLLSLQ
						CLISQIKMIMGAGWQ/SSGFTDE
						KETSKIKADPEKLKKLQESLLE
						EDSEEEGDLCRICQIAGGSPSNP
						LLEPCGCVGSLQFVHQECLKK
						WLKVKITSGADLGAVKTCEMC
						KQGLLVDLGDFNMIEFYQKHQ
						QSQIKTEATVVEHKDIYNVTET
						QLVPEARSVLLVGAGQMLLTG
						ANRALNSDYSSMRPKVKAQNE
						LMNSGLYLVLLLHLYEQRFAEL
						MRLNHNQVERERVRVTRVLGG
						LSPNVSCGVVSGVGSFQWVRG
						LTDLKNEASDLHAQIPAE
23091	53459	Α	23225	1	594	MIAAFNPGKVDTVTTNGPFIDL
						NYMVYMFHYDPTHGKFHDVP
						MFVMGMNHKHENKLTVISNAS
Ī						CTTNCLDPLAKVIHDNFGIREG
						LMTTVHSHATQKTVDGPSRKL
						WHDSHGAFQNTIPASTGAAK\I
						KKVVRQA/SQGPLKGILGYTEH
						QVVPLEFKSNTRPSTFEAGAGI
1						AVNDHFVKLISWYGNEFAYSN
						KVIKPM
23092	53460	Α	23226	3	232	
23093	53461	Α	23227	1078	2221	QQTLQPAALPCPFPPPTHPPEVL
		İ				PTAMNTTSSAAPPSLGVEFISLL
						AIILLSVALAVGLPGNSFVVWSI
						LKRMQKRSVTALMVLNLALAD
						LAVLLTAPFFLHFLAQGTWSFG
						LAGCRLCHYVCGVSMYASVLL
						ITAMSLDRSLAVARPFVSQKLR
						TKAMARRVLAGIWVLSFLLAT
						PVLAYRTVVPWKTNMSLCFPR
						YPSEGHRAFHLIFEAVTGFLLPF
						LAVVASYSDIGRRLQARRFRRS
						RRTGRLVVLIILTFAAFWLPYH
	1					VVNLAEAGRALAGQAAGLGLV
					1	GKRLSLARNVLIALAFLSSSVNP
		1				VL\YACAGGG\LLRSAGVGFF\A
						KLLEG\TGSEASSTP\RGGSLGQ
						TV\KSGPAALEPGPSESLTASSP
						LKL\NELN
23094	53462	A	23228	6125	6319	ASVVPVTSALAMCSPKSGIGDG
						FAVLRVEDVAQK***KAALGV
		<u> </u>				ASSPAWQQTREREHVQGNSPL
23095	53463	A	23229	1	1347	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
23096	53464	A	23230	1	1124	MRECISIHVGQAGVQIGNACWE
20070		1				LYCLEHGIQPDGQMPSDKTIGG
		İ				GDDSFNTFFSETGAGKHVPRAV
		1				FVDLEPTVVDEVRTGTYRQLFH
]		}				PEQLITGKEDAASNYARGHYTI
						GKEIVDLVLDRIRKL\FGGGTGS
						GFASLLMERLSVDYSKKSKLEF
						AIYP\APQVSTAVVEPYNSILTT
1						
						HTTLEHSDCAFMVDNEAIYDIC
						RRNLDIERPTYTNLNRLIGQIVS
						SITASLRFDGALNVDLTEFQTN
						LVPYPRIHFPLATYAPVISAEKA
		İ				YHEQLSVAEITNACFEPANQMV
						KCDPRHGKYMACCMLY\RGDV
						VPKDVNAAIATIKTKRTIQFVD
						WCPTGFKV*LGDMEAFQQAAD
		1				AQNNTGLEGPHPLGRLSLFHGL
23097	53465	Α	23231	1	2918	MDVYCAHESLRHTKSPEATGQ
	-					TSGQRPEALQAFWPAFGSSPVT
						GSSPVTEVTTKGRWSSSRQVL
						WDRIPLAKSRPAISNRQSQRTNI
						PWWESCPPILNLEREQPPRFAQP
						GTYASRWKALEEMEKQQREQ
						VDRNIREAKEKLEAEMEAARH
		1				EHQLMLMRQDPGLALVSCMQ
						QGQSMSRSPKKVFKHLYELNL
				ļ		TKLQSLGSRSRLRWKDSPFWGE
						EAGGKRPPLVSLALHRSPSLLL
						LFVPKHPSKGRPDCTVQRS
23098	53466	A	23232	1	158	
23099	53467	A	23233	1	414	LGGLKGTWGETKPAMAAAEEE
						DGGPEGPNRERGGAGATKTLL
				ţ		YTGNHLFSPQWLETRPERQECP
						VCKAGISREKVVGEKSLLTFSL
	1	1				HFLRLKTPQICRLPV*LFLPP*GF
						QPFGDTGGFHFSFGVGAFPFGF
		$oldsymbol{ol}}}}}}}}}}}}}}}}}$				FNSPSY
23100	53468	A	23234	1	663	
23101	53469	Α	23235	2	667	SWAGGGFWFVGACVCVFGGTE
						GYVGRNKTGHGSSRRRRSGAP
						NVPNRERGGAGRPKTPFECNIC
						LETARELVLSVLANLYFWTISH
						QV\LE\TRPE\RQECP\VCKAGISR
						EKVVP\LYGRGSQEAPRIPRLKT
				İ		PPPPPRGQR\PAPE\SRGGF\QPFG
						DTGGFHF\SFGVGAFPFGFFTTV
						F\NAHEPFRRGTGVDLG\QGHP
						A\SSW\QD\SLLLFLAIFFFF\WLL
23102	53470	Α	23236	363	476	
	53471	В	23237	224	364	T

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
23104	53472	A	23238	1	1324	MRQRRAELLSKDKRVRTDGNN
						KCSPSAIGKRKERGTNDWIEGD
						NNQESQGLNCSDTLNRDLGPN
						TRGFLYAGLSGLDPDPSLPTPD
						MSSEVLEDNLDTLSLYSGKDSD
						SVKLLEEYADSESQASLQGRSR
						CWQHPHPAPLSYPPAASGSQLS
						LQGKFPSAKPPLLGGYTCTAQR
						TQWALTKRSLRATRESVSRQQ
						GVYGLVGRKRTRHHRGVLGLL
						GIPSARPERPGTGSEAGPTRLPG
						AAVARRRPARPVRGRLGDPAP
						TAAQTSGPASLPPPPPPRSYGDP
						HQNGYPDPDPAPSASPDLPPAA
						GDTRGSGGRRKKRRTGTARLN
						TPGRKA\PKQKAAAGRKGRAH
						QPAPRTPEPQRRRPIRPPNQTDR
						LRHPHRRAGRGPPPGGAGGTA
						NPSTGANGGAATARRTAGART
	}	ļ				RGARGASERAGRHGSRPASWL
		L				TATPPPGT
23105	53473	Α	23239	1	433	PTRPSSPEETDGPWDPEGHPGG
		1				PLTDPPNAPVLALPPAIPGLSLS
		1				MADGSSDAAREPRPAPAPIRRR
						SSNYRAYATELQDLCRQLHAR VDKVDEERYDIEAK\VTKNITEI
						ADLTQKIFDLRGKFKRPTLRRV
						RISADAMMQALL
23106	53474	A	23240	2	409	RISADAMINQALL
23107	53475	A	23240	3	490	SSPEETVGPWDPEGHPGGPLTD
23107	33473		23241	-	1470	PPNAPVLALPPAIPGLSLIMADG
						SSDAVRAAG*GVAGTPRARAR
						EPRPAPAPIRRRSSNYRAYATEP
						HAKKKSKISASRKLQLKTLLLQ
Ì						IAKQELEREAEERRGEKGRALS
						TRCQPLELAGLGFAELQDLCRQ
						LHARVDK
23108	53476	Α	23242	149	846	VSAWRNGSSDAAREPRPAPAPI
						RRRSSNYRAYATEPHAKKKSKI
						S\ASRKLQLKTLLLQIAKQE\LE
						REAEERRGREGARSLAPRC\QP\
						LELAG\LGFADLPGLCCRQFHA
						RV\DKVDEERYDIEAKVSKNITE
-						IAN\LT\QKIFDLRSKFKRPTL\RR
1						VRISADAMMQALLGARAKESL
						DLRAHLKQVKKEDPEKENREV
						GDWRKNIDALSGMEGRMKKW
						RELSLPAYCPCPEEGH
23109	53477	Α	23243	1	487	
23110	53478	Α	23244	80	511	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		1		sequence		
23111	53479	A	23245	427	1417	I KTMWKVEFGVRFSITRLLLEHQ
23111	33479	^	23243	427	1417	PKQTMYIPLPK/PPLWPEEPKTE
						-
						APKQTPRAPPKPKTSPRPRIPQT
						QPEETDQSTQEPFTTKIPRTTEL
						AKTTQGLEMLSCLIHNPKMHK
						DSTNEQIAASLQACYYEQLLM
						YIRVTDVDSVGPGNGKEAVTA
						PQAPTLFWLNHKAQLAKDWA
						GKDVNITPTFATSKKENKPPNN
						LQICNVSKIHQRAKVAGKPEIK
						DKHIQKEMPLDAIEASKINLVKI
						FNEFLKAWKCRLSFEYRIPENQ
						GHKGSFYLLVGSSRTSWLLTEK
						IGSSKRVLRMPPMDILKTTQILL
						SFLWSINLTPRGKNNHLVAPAH
23112	53480	A	23246	3	1011	
23113	53481	Α	23247	2	334	PIQSFFLVLKKPPLRKGVYLSSK
			1			SLSRWFFKVQAN*RSQPQKAFS
						RLSCHC
23114	53482	A	23248	1	642	
23115	53483	A	23249	854	1150	KEDTNKWKNIPCSWLGRINIVK
		ŀ				MAIMPKVNL*IQCHPHQATNDF
					ŀ	SSQNWKKTTLKFIWEPKKEPAL
						PSQS*AKRAKLGGITLPDFQTIT
						YKATVTKTA
23116	53484	A	23250	1754	2393	PSNRREFGESWPETQAPGALRP
		İ				LLPAPGTPRARERPAGWPARPM
Ì						EVYIPSFRYEESDLERGYTVFKI
ł						EVLMNGRKHFVEKRYSEFHAL
1		1				HKKLKKCIKTPEIPSKHVRNWV
		1			İ	PKVLEQRRQGLETYLQ\AVILE
						NEELPKLFLDFLNVRHLPSLPK
						AESCGSFDETESEESSKLSHQPV
						LLFLRDPYVLPAASDFPNVVIE
						GVLHGIFYPHLQPR
23117	53485	A	23251	1	265	AGNRLRFQLELEFVQCLANPNY
23117	22702		23231		203	LN/WYPQCLHMLELLQYEHFR
						KELVNAQCAKFIDEQQILHWQ
						HYSRKRMRLQQAL\AKQQQQN
		1				NTSGK
23118	53486	A	23252	1	987	IVIOUR
23119	53487	A	23252	1	491	RGFRNFARVSGLLLCQAGGVL
واالوع	101	``		1		VSSFVMAAAVAMETDDAGNR
						LRFQLELEFVQCYANHNTLIFLP
		1				QR\GYF\KDKAFVNYLKYLLYW
						\KDPEYAKYLKYPQCLHMLGA
						•
						APNMEHFRKEAGWNAQC\AKF
						IGWNRQVSTLGKHYS\RKRM\R
			L	1		LQQALAEQQQQNNTSGK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23120	53488	Α	23254	73	355	HEGQVHVLLASAHVDVAIVLG
						AGSMVEGEARPEQAAQHHQGI
		l				APLHLDAPGPSWTQSRSPRNRA
						NQNHTCSEAT*SHPEQLPQTLW
						ECHAPRSP
23121	53489	В	23255	1	441	
23122	53490	A	23256	528	698	HGQRFLHGPV/H/HHSHYDN/PH
		ĺ				HSLHHHRCLHHCHLHHHRHHL
		ļ				HHHPLHCYHHHPGQNG
23123	53491	Α	23257	1	403	
23124	53492	Α	23258	509	4547	LYLTTSRREEHLKSLVVETQDL
						ASPVLRSVSICTKVEEAFRQAH
		l		-		VIVVLDDSTNKEVFTLEDCLRS
				-		RVPLCRLYGYLIEKNAHESVRV
		l				IVGGRTFVNLKTVLLMRYAPRI
						AHNIIAVALGVEGEAKAILARK
		1				LKTAPSYIKDVIIWGNISGNNYV
İ				,		DLRKTRVYRYESAIWGPLHYSR
		i				PVLNLIFDSEWVKREFVAILKN
						LTTTGRQFGGILAAHSIATTLKY
					:	WYHGSPPGEIVSLGILSEGQFGI
						PKGIVFSMPV
23125	53493	Α	23259	56	290	
23126	53494	Α	23260	468	993	VLLACFWPGSDFWPRSRRKTY
	1					GTAPQSCYSFYVALDCGGVSRL
	Ì					VIFVSWRNPQVAPTSAHQNRPS
						RNPVSRPPNTQRVARRKHYAL
						ADGYTERRWTNAPCRAESSFPP
						NCPSAAPIPDSYDK*PSRATLFT
						SHLTVEAFPVWSYLYPGETPRL
						HPRAHTRTVPAAIRYPDRQTPN
						GLRGANTMHSQMDTLNGDGR
						THLAGQRARSPRTARPQRQYRI
						HTTSDLPGLVRKYASGCCLRIW
						YCCAWIERALSLQVGGGVFGD
	<u></u>					PLSEGQAECDHGST
23127	53495	Α	23261	1	3114	TOOL OL OOKOODYWYDY DDOYG
23128	53496	Α	23262	3	401	TSSLSLSGKSGRYIVFLRRSVGI
						QSPSAVATVRLLLAGSDRRFAA
						GSAGCAVLSRAERS*EPGCCYR
						IRRSAFRLRAGSIGNRDKPVAW
						PDTGSGDNTEVYFRAHQAGILP
00100	52.405	<u> </u>	222.62	0.074	20/2	DTVLAGPAQRVAVSGGTPVDW
23129	53497	Α	23263	2874	3062	

SEQ ID	SEQ ID NO:	1	-	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23130	53498	Α	23264	2766	3323	MPDLVEGREKSDKGAQPQPRT
						QSRSARTNGSGVRILHSLWRIIG
		ŀ				IPLLLGYSLVCSRVLLACFWPG
						SDFWPRSRRKTYGTAPQSCYSF
						YVALDCGGVSRLVIFVSWRNP
						QVAPTSAHQNRPSRNPVSRPPN
						TQRVARRKHYALADGYTERR
						WTDAPCRAESSFPPNCPL*RQY
						RIHIVDASVCGP
23131	53499	Α	23265	968	1440	
23132	53500	Α	23266	3	640	LSVVRSRIHMGVGVGVGCPVR
			Ì			YRNQDDHELQITHGNKILCGIV
						CDKGAQPQPRTQSRSARTNGSG
			:			VRILHSLWRIIGIPLLLGYSLVCS
						RVLLACFWPGSDFWPRSRRKT
						YGTVPQSCYSFYVALDCGGVS
ŀ						RLVIFVSWRNPQVAPTSAHQNR
						PRRNPVSRPPNTQRVARRKHYA
						P\ADGYTERRWTIAPCRAESSFP
		ļ				PNCPSAAPIPDSYDK
23133	53501	Α	23267	1	3140	MVYKMRYRSQHPYSIKEKQMK
						SEVLSVKEKIGYGMGDAASHII
						FDNVMLYMMFFYTDIFGIPAGF
						VGTMFLVARALDAISDPCMGL
						LADRTRSRWGKFRPWVLFGAL
						PFGIVCVLAYSTPDLSMNGKMI
]	YAAITYTLLTLLYTVVNIPYCA
						LGGVITNDPTQRISLQSWRFVL
						ATAGGMLSTVLMMPLVNLIGG
						DNKPLGFQGDLSPMFSTPEEIA
						RPGPYENDVHVVGASSLAAGH
		ļ	22262			KTLIPELVRSAEQHMGTR
23134	53502	Α	23268	750	1340	ILHAPAPPFSASASHEQPEWSD
						KGAQPQPRTQSRSARTNGSGVR
						ILHSLWRIIGIPLLLGYSLVCSRV
						LLACFWPGSDFWPRSRRKTYG
	Ī			1		TAPQSCYSFYVALDCGGVSRLV
						IFVSWRNPQVAPTSAHQNRPSR
						NPVSRPPNTQRVARRKHYALA
						DGYTERRWTNAPCRAESSFPPN
22125	50500	ļ. —	222.60		600	CPFAAPIPDSYDK*PFRIVS
23135	53503	Α	23269	93	699	RASVQQKKLSTDDKGAQPQPR
				1		TQSRSARTNGSGVRILHSLWRII
		1				GIPLLLGYSLVCSRVLLACFWP
						GSDFWPRSRRKTYGTAPQSCYS
		İ				FYVALDCGGVSR\RSYLYPGET
						PRVAPTSAHQNRPSRKPG\TRPP
						NTQRVARRKHYALADGYTER\
						SATNAPCRAESSFPPELPVPQVP
		1				NTGFIRQVTFRDCFVNTPQAAA
		<u></u>				SGFW

SEQ ID			SEQ ID NO:	4	•	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
			- ',	sequence		
22125	[50504		100070	1.000	11500	lonvon vinn vin von on on on on
23136	53504	Α	23270	1000	1530	QRYSDNPPNDKGAQPQPRTQSR
		1				SARTNGSGVRILHSLWRIIGIPL
		l				LLGYSLVCSRVLLACFWPGSDF
						WPRSRRKTYGTAPQSCYSFYV
						ALDCGGVSRLVIFVSWRNPQV
						APTSAHQNRPSRNPVSRPPNTQ
						RVARRKHYALADGYTERRWT
22125		ļ. —	22271	0.10	711	NAPCRA/VEPFPPNCPWEAQYRI
23137	53505	A	23271	242	711	LKQKRDKEDRSSRVSAVLLIK/
ļ		1				HRRPLPLPIGLKACHCSCTAKC
ţ						LGGCESPRKGGQLLIFEGHVKI
ł						DLEIFTKSLWLTRQLIVKKETEK
						AVKHASNICIMRTGLTFQSHKIE
	l	ŀ				DGLFGLLVIATEAEWSYVCQST
		<u> </u>				NPVDVYPLTAVPACCIISASDVP
23138	53506	Α	23272	2	459	CVNSQIGRGHKITTYPFSPRKSG
	-	1				RKGMAQSQGWVKRYIKAFCK
	}	ł				GFFVAVPVAVTFLDRVACVAR
						VEGASMQPSLNPGGSQSSDVVL
İ		i				LNHWKVRNFEVHRGDIVSLVS
		l	İ			AKNPEQKIIKRVIALEGDIVRFP
		ļ				*DFCMPMPHISCGPQSAGRNW
23139	53507	A	23273	1	474	
23140	53508	Α	23274	358	948	PLTPFSPRKSGRKGMAQSQGW
ļ		1				VKRYIKAFCKGFFVAVPVAVTL
						LDRVACVARVEGASMQPSLNP
					1	GGEPV/ASDVVLLNHWKVTEF/
		1			ļ	SEVHRGDIVFIG\APKNPEQKIIK
		l				RVICSLKGDIFQNHRTQKP\DVK
		1				VPRGSHLGLEGDHSWDTVFDS
		1				NSFGPVSLGLLHAHATHILWPP
						ERWQKLESVLPPERLPGTERRG
	53.500	<u> </u>	22275			MTA
23141	53509	A	23275	90	664	NKYRLQPSPPRLALWDRASDPP
		ĺ				APPEDPETAPAMEEDQELERKI
		l				SGLKT\SMVEG\ERKTT\LELV\Q
						AAG\TDRP\CVTFVLHEEDHTLG
						\NFLRY\MIMKNP/EKVEFCGYT
1						YG/HPSFQRAKLNLPHSRFRGYP
						FPAVEPISREALNELM\NVC\QP
						CAFTSFRASIKDYKESKSKQEM
						ESTFLVLLWQFYKGGTVL
23142	53510	С	23276	108	320	
23143	53511	В	23277	446	625	
23144	53512	Α	23278	1	3042	
23145	53513	Α	23279	1	3156	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
•	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23146	53514	Α	23280	518	2076	QQRR*MCQNLMTH\QNQRSGR
23140	122214		23200	310	2070	SQNDF*RRRKNISERSAVYR*R
ŀ						QLYIRRQPYAERPS\HVEDKGH
ľ						
ł						KYLVFEANTGTENGYQGEESLF
						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTLKKVMKPLITSNTVTDEIER
		j				ANVFKMNGKWYLFTDSRGSK
				1		MTIDGINSNDIYMLGYVSNSLT
1						GPYKPLNKTGLVLQMGLDPND
						VTFTYSHFAVPQAKGNNVVITS
						YMTNRGFFEDKKATFAPSFLM
İ			İ			NIKGNKTSVVKNSILEQGQLTW
1						LQVAKRAGLGGGQSGRTVLRE
				1		RVRIEIASTHIALAVECVSVRV
ł				_		WKVPRLPSRQKYAKHASQLVS
Ì						NQKSEEEEYMKTFNISQQDLEL
}						VEVATEKITMLYEDNKHHVGA
						AIRTKTGEIISAVHIEAYIGRVTV
			-			CAEAIAIGSAVSNGQKDFDTIV
ŀ						AVRHPYSDEVDRSIRVIPTYGT
						DEWEQWWNAFNEENLFCSEE
						MPSSDDEATADSQHSTPPKKKR
						KYIYVLVIAITMVMRFWQYING
23147	53515	Α	23281	299	457	
23148	53516	Α	23282	584	1185	QQRR*MCQNLMTH\QNQRSGR
					i	SQNDF*RRRKNISERSAVYR*R
ŀ		ļ				QLYIRRQPYAERPS\HVEDKGH
						KYLVFEANTGTENGYQGEESLF
						NKAYYGGGTNFFRKESQKLQQ
		l				SAKKRDAELANGALGIIELNND
ŀ						YTSEKSNEAADHFKHVPQAKG
						NNVVITSYMTNRGFFEDKKATF
		İ				APSFLMNIKGNKTSVVKNSILE
						QGQLTVN
23149	53517	A	23283	1	3618	AOADIAIA
23149	53517	A	23284	191	819	VNVTSFGSRPICSTSPVLFSGL*
23130	0100		23204	171	017	GPVKEFDT*PSM*ISFELIPSIVIF
						EPRESVNKYHLPFILKTFARSISS
1					,	
						VTVFEVISGFITFFNV*SLFNSMI
İ						PRAPFANSASRFLALC*SFWLSL
						RKKFVPPP*YALLNKDSSPW/SS
]						VFCSRVGFEYKVFMAFVFNVV
						RVSQRMVVAGCIIAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23151	53519	A	23285	2	2556	WRKIYQAN/GKTKKAGVAILVS
				_		DKTDFKPTKIKRDKEGHYIMVK
						GSIQQEELTILNIYAPNTGAPRFI
						KQVLSDLQRDLDSYTIIMGDFN
						TPLSTLDRSMRQKVNKNTQEL
						NSALHQADLIDIYRTLHPKSTEC
		İ				TFFSAPHHMYSKIDHIVGCKAL
						LSKFKRTEIITNCLSDHSAIKLEL
						RIKKLTQNCSTTWKLNNLLLND
						YWVHNEMKAEIKMFFETNENK
		l				DTTYQNLWDTFKAVCRGKFIA
						LNAHKRKQERSKIDTLTSQLKE
			•			
						LEKQEQTHSKASRRQEITKVRA
						ELKEIETQKTLQKINEFRSWFFE
						RINTIDRPLARLIKKKREKNQID
						AIKNDKGDITTDPTETQTTIREY
						YKHLYANKLENLEEMDKFLDT
						YTLPRLNQEEVESLNRPITGSGI
	<u> </u>					VAIINSLPTKKSPGPDGFTAEFY
						QRYKEELVPFLLKLFQSIEEEGI
						LPNSFYEASIILIPKLGRDIAKKE
						NFRPISLMNIDAKILNKILAHRI
						QQHIKKLIHHDQVGFIPGMQG
						WFDIRKSINVIQHINRTKDKNH
		l				MIISIDAEKAFDKIQQPFMLKTL
		İ				NKLVLEVLVRAIRQEKEIKCIQL
		ł				GKEEVKLSLFADDMIVYVENPI
		1				VSAQNLLKLISNFSKVSGYKIN
						VQKSQAFLYTNNRQTESQIMSE
						LPFTIASKSIKYLGIQLTRHVKD
						LFKDNYKPALNEIKEDTNKWK
		1				NIPCSWIGRINIVKELEKTTLNFI
23152	53520	Α	23286	1	3087	
23153	53521	Α	23287	1	741	MLGKKKPVCEVGGDSGVPKKE
						FQEGRQVLFKLWRGQANKVG
						DNSIDSWKNAGRVFKDSDKFD
						ANDPILKDQTQEWSGSATFTSD
						GKIRLFYTDYSGKHYGKQSLTT
		l				AQVKVSKS/V*HTQNQRSGRSQ
					i	NDF*RRRKNIS*RSAVYR*RQL
						YIRRQPYAERPS\HVEDKGHKY
						LVFEANTGTENGYQGEESLFNK
						AYYGGGTNFFRKESQKLQQSA
						KKRDAELANGALGIIELNNDYT
						LKKVVKPLITSN
	L		L		<u> </u>	LIXIX V VIXI LI I DIV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence -		
23154	53522	Α	23288	849	1688	QQRR*MCQNLMTH\QNQPSGR
						SQNDF*RRRKNISERSAVYR*R
						QLYIRRQPYAERPS\HVEDKGH
						KYLVFEANTGTENGYQGEESLF
		İ				NKAYYGGGTNFFRKESQKLQQ
•		İ				SAKKRDAELANGALGIIELNND
İ					ĺ	YTLKKVMKPLITSNTVTDEIER
						ANVFKMNGKWYLFTDSRGSK
						MTIDGINSNDIYMLGYVSNSLT
						GPYKPLNKTGLVLQMGLDPND
		ŀ				VTFTYSHFAVPQAKGNNVVITS
		1				YMTNRGFFEDKKATFAPSFLM
		1				NIKGNKTSVVKNSILEPGQLAV
23155	53523	Α	23289	1	3171	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRIEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKTEIKMFF
						ETNENKDTTYQNLWDAFKAVC
				1		RGKFIALNAYKRKEERSKIDTL
		1				TSQLKELEKQEQRHSKPSRRQE
		İ				ITKMRAELKEIETQ
23156	53524	Α	23290	1	3255	
23157	53525	A	23291	1	3924	
23158	53526	A	23292	1	3139	
23159	53527	Α	23293	1	3210	MVKGSIQQEELTILNIYAPNTG
				:		ALRFIKQVLRDLQRDLDSHTIIM
						GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
						ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
		I				FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQ
23160	53528	В	23294	1	3127	
23161	53529	Α	23295	1	3325	
23162	53530	Α	23296	1	3145	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23163	53531	ĪA	23297	191	819	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
		1		ļ		EPRESVNKYHLPFILKTFARSISS
	ļ	ŀ				VTVFEVISGFITFFNV*SLFNSMI
						PRAPFANSASRFLALC*SFWLSL
Ì		1				RKKFVPPP*YALLNKDSSPW/SS
						VFCSRVGFEYKVFMAFVFNVV
						RVSQRMVVAGCIIAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23164	53532	A	23298	1	3352	
23165	53533	A	23299	1841	2679	QQRR*MCQNLMTH\QNQRSGR
		1				SQNDF*RRRKNISERSAVYR*R
l l						QLYIRRQPYAERPS\HVEDKGH
					4.	KYLVFEANTGTENGYQGEESLF
ĺ						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTLKKVMKPLITSNTVTDEIER
			:			ANVFKMNGKWYLFTDSRGSK
	1					MTIDGINSNDIYMLGYVSNSLT
						GPYKPLNKTGLVLQMGLDPND
						VTFTYSHFAVPQAKGNNVVITS
						YMTNRGFFEDKKATFAPSFLM
						NIKGNKTSVVKNSILEQGQLTV
23166	53534	Α	23300	1207	1533	YVHCKPSGWTVRTFDKPRKRFI
						AFFIAGILFRAIKNHFLPRETLQ
						CLPYILTGFRRGQSEYFSIFSNM
						DLADTVMFL*GAIRFSDLVNEQ
		1				IQHTFLIPGETICRLSEVV
23167	53535	Α	23301	1	3253	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQGDLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
		İ				GSKALLSKWKRTEIITNYLSDH
						SAIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
ļ						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
						KNQIDTIKNDK
23168	53536	Α	23302	1	3095	
23169	53537	Α	23303	2	2678	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \= possible nucleotide insertion)
				sequence		
23170	53538	Α	23304	1	3335	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
1						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
•						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
		}				RGKFIALNVYKRKQERSKIDTL
ļ						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQ
23171	53539	В	23305	1	2293	THURSTEDING
23172	53540	Α	23306	1	3810	
23173	53541	A	23307	255	1215	CVGQLAAAKR*RNSS*IQRLSR
						CVCSCGKPERR**HINLHVLSK
				•		GRRQLN/QTAGKTRAVSLKTAI
						SSTPTIRS*KIRRKNGPVLQPLH
						LTEKSVYSTLTIPTKAINTLYSK
						PTREQKTDTKAKNLYLTKRTT
		l				AAARTSSVKKARSFSRALKNA
		ŀ				MLS*RTAPSVS*S*IMITH*KK**
			:			SR/CITSNTVTDEIERANVFKMN
						GKWYLFTDSRGSKMTIDGINSN
			ŀ			DINMLGYVSNSLTGPYKPLNKT
		İ				GLVLQMGLDPNDVTLTNSHFA
						VPQAKGNNVVITSYMTNRGFF
						EDKKATFAPSFLMNIKGNKTSV
]				VKNSILEQGQLTVN
23174	53542	Α	23308	1	4539	
23175	53543	A	23309	1	3229	
23176	53544	В	23310	1	3552	
23177	53545	A	23311	1	3139	
23178	53546	Α	23312	1	3057	NGW ON TO WITHOUT A GODD
23179	53547	Α	23313	1	3215	MGKKQNRKTGNSKTQSASPPP
1		1				KERSSSPATEQSWMENDFDELR
		1				EEGFRRSNYSELREDIQTKGKE
						VENFEKNLEECITRITNTEKCLK
		1				ELMELKTKARELREECRSLRSR
1						CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
]			KRPNLRLIGVPESDVENGTKLE
]			NTLRDIIQENFPNLARQANVQIQ
						EIQRTPQRYSSRRATPRHIIVRFT
						KVEMKEKMLRAAREKDRSTRQ
		1	L			KVNKDTQELNSA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23180	53548	Α	23314	1	1921	MVKGSIQQEELTILNTYAAHTG
						APRLIKQVLSDLQRDLDSHTIIM
						GDFNTPLSTLDRSTRQKVNKDT
						QELKSALHQADLTDIYRTLHHK
						STEYTFFSAPHHIYSKIDHILGSK
						ALLSKCKRTEIITNYLSDHSAIK
						LELWIKNLTQNHSTTWELNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYHNLWDTFKAVCRGK
						FIPLNAHKRKQERSKIDTLTSQL
1						KELEKQEQTN\HSKASRRQEITK
						IRAELKEIETQKTL*KINESRSW
						FFERINKIDRLLARLIKKKREKN
						QIDAIKNDKGDITTDPTEIQTTIR
						EYCKHLYANKLENLEEMDKFL
						DTYTLPRLNQEEVESLNRPITG
		ļ				AEIVAIINSLPTKKSPGPDGFTA
						KFYQRYKEELVPFLLKLFQSIEK
						EGILPNSFYEASIILIPKPGRDTT
						KKENFRPISLMNIDAKILNKKL
						AKRIQQHIKKLIHHDQVGFIPG
}						MQGWFNIRKSINVIQHINRAKD
ľ						KNHMIISIDAEKAFDKIQQPFML
						KTLNKL\GIKYLGIHLTRDVKDL
		1				FKENYKPLLKEIKEDTNKWKNI
		1				PCSWVGRINIVKMAILPKMCL*
		į				RRWTPLCLLTAAASLNRE*ERL
						VQPLLWRQMYCGPRHCRQVPQ
		<u> </u>			,	HRRLNWSPSLRLS
23181	53549	В	23315	1	2907	OODD*MCON! MTUVONODCCD
23182	53550	Α	23316	584	1278	QQRR*MCQNLMTH\QNQRSGR
						SQNDF*RRRKNISERSAVYR*R
						QLYIRRQPYAERPS\HVEDKGH
						KYLVFEANTGTENGYQGEESLF
						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTLKKVMKPLITSNTVTDEIER
						ANVFKMNGKWYLFTDSRGSK
						MTIDVPQAKGNNVVITSYMTN
1						RGFFEDKKATFAPSFLMNIKGN
02192	62551	_	22217	1	2245	KTSVVKNSILEQGQLTVN
23183	53551	Α_	23317	1	3345	

SEQ ID			SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23184	53552	A	23318	2	3366	WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMVK
						GSIQQEELTILNIYAPNTGAPTFI
						QQVLSDLQRDLDSHTLIIGDFN
						TPLSTSDRSTRQKVNKDTQELN
						SALHQADLIDIYRTLHPKSTEYT
		1				FFSAPHHTYSKIDHILGSKALLS
						KCKRTEIITNYLSDHSAIKLELM
						IKNLTQNHSTIWKLNNLLLNDY
						WVHKEMKAEIKMFFETNENKD
			•			TT\YQNLWDTFKAVCRGKFIAL
		1				NAHKRK\QER
23185	53553	A	23319	1	2611	MGDFNTPLSTLDRSTRQKANK
						DTQELNSALHQVDLIDIYRTLH
						PKSTEYTFFSAPHHTYSTTDHIL
						GSKALLSKCRRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYGVHNEMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVC
				İ		RGKFIALNAHKRKQERSKIDTL
		ļ				TSQLKELEKQEQTHSKASRRQE
						ITKTRAELKEIETQKTLQK\INES
			ŀ			RSWFFERINKIDRPLARLIKKKR
		1				EKNQIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DKFLHTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLPTKKSPGPDG
		ĺ				FTAEFDQRYKEELVPFLLKLFQ
						SIEKEGILPNSFYEASIILIRKPGR
		1				DTTKKENFRPISLMNTDAKILN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWFNIHKSINVIQHINRA
						KDKNHMIISIDAEKAFDKIQQPF
		İ				MLKTLSKLGIDGTYFKIIRAIYD
					1	KPTANIILNGQKLEAFPLKTGTR
		l				QGCPLSPLLFNIVLEVLARAIRQ
						EKEIKLISNFSKVSGYKINVQKS
						QAFLYTNNRQTESQIMSQLPFTI
						ASKRIKYLGIQLTRDVKDLFKE
						NYKPLLKEIKEDTNKWKNIPCS
						WVGRINIMKMAILPKVIYRFNA
						IPIKLPMTFFTELEKTTFKFIWN
						QKRARITKSILSQKNKAGGITLP
L	<u> </u>	<u>L</u> _				DFKLYYKATVTKTAWYWYQN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		į		sequence		
23186	53554	A	23320	11	3099	MGELITPLSTLDRSTRQKVNKD
						TQELNSALHQGDLIDIYRTLHP
		ļ				KSTEYTFFSAPHHTYSKIDHILG
						SKALLSKCKRTEIITNYLSDHSA
						IKLELRIKNLTQNRSTTWKLNN
						LLLNDYWIHNEMKAEIKMFFET
						NENKDTTYQNLWDAFKAVCR
		Ì				GKFIALNAHKRKQERSKIDTLT
		İ				SQLKELEKQEQTHSKASRRQEI
1		1				TKIRAELKEIETQKTLQKINESR
		l				SWFFERINKIDRPLARLIKKKRE
						KNQIDTIKNDK
23187	53555	A	23321	911	1539	VNVTSFGSRPICSTSPVLFSGL*
25107	33333	 	25521		1337	GPVKEFDT*PSM*ISFELIPSIVIF
		ļ				EPRESVNKYHLPFILKTFARSISS
	}					VTVFEVISGFITFFNV*SLFNSMI
		İ				PRAPFANSASRFLALC*SFWLSL
		ŀ	Ì			RKKFVPPP*YALLNKDSSPW/SS
		ŀ				VFCSRVGFEYKVFMAFVFNVV
]				RVSQRMVVAGCIIAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23188	53556	Α	23322	191	820	VNVTSFGSRPICSTSPVLFSGL*
25100	22230	^	23322		020	GPVKEFDT*PSM*ISFELIPSIVIF
		ł				EPRESVNKYHLPFILKTFARSISS
		i				VTVFEVISGFITFFNV*SLFNSMI
		1		!		PRAPFANSASRFLALC*SFWLSL
						RKKFVPPP*YALLNKDSSPW/SS
						VFCSRVGFEYKVFMAFVFNVV
		1				RVSQRMVVAGCIIAFIDKLLNV
		l				LICFSVSVKNRFVIFHSVDFECV
23189	53557	A	23323	965	5050	TWKGTTSTSRCKIMPKYRSTRQ
23109	33337	<u> </u> ^	23323	903	3030	KVNKDTQELNSALHQADLIDIY
		1				RTLHTKSTEY/TFFSAPHHTYSK
						IDHIVGSKALLSKCKRTEIITNY
		1				LSDHSAIKLELRIKNLNQSRSTT
		l				WKLNNLLLNDYWVHNEMKAE
ŀ						IKMFFETK\ENK\DTTYQNLWD
]		ļ				AFKAVCRGKFIALNAHKRKQE
		1				RSKIDTLTSQLKELEKQEQTHS
						KASRRQEITKIRAELKEIETQKT
		1				LQKINESRSWFFERINKIDRPLA
22122	50.550	<u> </u>	10000 :	1	0707	RLIKKKREENQ
23190	53558	A	23324	1	2787	
23191	53559	Α	23325	1	902	
23192	53560	A	23326	1	3640	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23193	53561	Α	23327	1	3686	MHIVVETALSASWQNKAKPPA
						RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
						WTENDYDKLREEGFRRSNYSE
			,			LQEDIQTKGKDVENFEKNLEEC
						ITRITNTQKCLKELMELKTKAR
		1				ELREECRSLRSRCDQLEERVSV
						MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIQENF
Ì						PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
23194	53562	A	23328	1	3433	MGDFNTPLSTLDRSTRQKVNK
	33302	ľ.	25520	1		DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIITNYLSDHS
	:					AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
						KNQIDAIKNDK
23195	53563	Α	23329	1	3070	MEGEMNEMKREGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
	j		<u> </u>			PESDGENGTKLENTLQDIIQENF
						PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
						LRAAREKDRSTRQKVNKDTQE
						LNSALHQADLIDIYRTLHLKSTE
						YTFFSAPHHTYSKIDHILGSKAL
						LSKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQNRSTTWKLNNLLLN
1						DYWVHNKMKAEIKMFFETNEN
						\KDTTYQNLWDA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	ı	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23196	53564	A	23330	3	2372	TTNCLSDHSAIKLELRIKKLTEN
23190	33304	^	23330]	2372	HTTTWKLNNLLLNDYWVDNE
						MKAEIKMFFEINEKK\DTTSQNL WDTFKAVCRGKLIALNAHKRK
						1
						QERSKIDTLTSQLKK\LEKQEQR
						NSKASRKQEITKIRAELKEIETQ
						KKTLQKINESRSWFFEKINKIDG
İ						PLARLIKKKREKNQVDAIKNDK
						GDIITDPTEMQTTIREYYKYLY
	ļ					ANKLENLEEMDKFLNTYTLQR
						LNQEGIESLNRPITGSEIEAIINSL
		l	!			PTKKSPGPDGFTAKFYQRYKEE
	ŀ	l				LVYPEPRSESECLSNIREFLRGC
		ŀ				GASLRLESIREDRNGRSQKTVH
		l				TEGDMNMNIKKIVKQATVLTF
		l				TTALLAGGATQAFAKENNQKA
		1				YKETYGVSHITRHDMLQIPKQQ
						QNEKYQVPQFDQSTIKNIESAK
						GLDVWDSWPLQNADGTVAEY
]						NGYHVVFALAGSPKDADDTSI
1		l				YMFYQKVGDNSIDSWKNAGR
		l				VFKDSDKFDANDPILKDQTQE
						WSGSATFTSDGKIRLFYTDYSG
1						KHYGKQSLTTAQVNVSKSDDT
		1				LKINGVEDHKTIFDGDGKTYQN
		1				VQQFIDEGNYTSGDNHTLRDPH
						YVEDKGHKYLVFEANTGTENG
		1				YQGEESLFNKAYYGGGTNFFR
		1				KESQKLQQSAKKRDAELANGA
		ł				LGIIELNNDYTLKKVMKPLITSN
		l				TVTDEIERANVFKMNGKWYLF
22107	50565	١.	00001		2007	TDSRGSKMTIDGINSNDIYMLG
23197	53565	Α	23331	2	2987	APRFIKQVLSDLQRDLDSHTIIM
		1				GDFNTPLSTLDRSTRQKVHKDT
		1	ŀ			QELNSALHQADLIDIYRTLHPK
		1	ļ		}	STEYTCFSAPHHTYSKIDHIVGS
						KALLSKCKRSEIITNCLSDHSAI
					1	KLELRIKKLTQNRSTTWKLNNL
			r I			LLNDYWVHNEMKAEIRMFFET
						NENKDTTYQNLWDTFKAVCRG
}						KFIALNAHKRKQERSKIDTLTS
						QLKELEKQEQTHSKASRRQEIT
						KIRAELKEIETQKTL\QKINE\SR
00101	50555	ļ.—	00000	222	701	SWFF\ERINK
23198	53566	Α	23332	339	701	PDKEGNEIWVDMYTVKPSGWT
						VRTFDKPRKRFIAFFIAGILFRAI
						KNHFLPRETLQCLPYILTGFRRG
						QSEYFSIFSNMDLADTVMFL*G
				1		AIRFSDLVNEQIQHTFLIPGETIC
	20.55	<u> </u>	2222	1.00	500	RLSEVV
23199	53567	Α	23333	190	502	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23200	53568	Α	23334	4472	8032	KKSGNNRLMRVHGKKRFLWQ
		1				KQGVCEVEIQAQRESARDGTD
		l		ļ.		REDAESPHAEVTVPRGTRTEILT
1						LWASRDRPASACGVTVPKGST
1						THPGGMDRVCGVDTAVLHWM
						KQVQGISNLEVETGPRRRDMD
						EAGSRHSQQTNIGTENQTPHVL
						THKWQLNRNSACDKLHKPHSV
						PGGVAAGDVGGKTISNPGVCAI
						EYQVDGSPHIWEDVNKGVYIEE
						DRAPLMERSSSPAMEQSWMEN
						DFDELREEGFRRSNYSELRED
23201	53569	Α	23335	1	2374	MVSISDLVICPPRHPKVLGLQGP
						PGLDSISDPSAGAGFLDWGEIG
						MPGPGRAGHQALCKCDCQCLE
						KTTTKAPGKMPKSTRSGPVRV
						RLADGPNRCAGRLECGMPDAG
						EQCVMTTGTSGRHCGLLGTGL
						WKGYTDLTIIPPGPGTPPQERTC
						QGDYHSGGTWTHSPLETTRRP
					<u> </u>	GSSSPAIRRLPAQMLLLPARPPH
						PRSSSPEAMDPPPPKAPPFPKAE
						GPSSTPSSAAGPRPPRLGRHLLI
		İ				DAN/GVYPYTYTVQLEEEPRGP
						PQREAPPGEPGPRKGYSCPECA
						RVFASPLRLQSHRVSHSDLKPF
						TCGACGKAFKRSSHLSRHRATH
		İ				RARAGPPHTCPLCPRRFQDAAE
						LAQHSWGTPRGPLLAAACNCE
						VARGRLESPGPERLLHGYGGRE
						EEGGWGRAAGGLDRVEGFISS
						KAHHYLLIDTQGVPYTVLVTRS
		1				HRGSQGPVGLQARKVLQLPRV
}]	LKGLRVHVHLQRHSITHSEVPQ
1						DFAGSLDSFQTPGESLRLVFRA
						LDTTQSSRISKAEPCLKEEPLSL
				1		GDLPYMHTTLCFCRKRRASPGP
}						GTLQRGALAWPDWASPRALPV
				1		PSLSSTTRSPAAPLFAVPLSGRT
						TQAMAFDGIIFQGQSQRSAGLT
				1		TTSRFLACQRPLRLCAWWASRS
						PRCTLRRPVGLRPGVHPRPRLV
						YRDLKPENVMASGQPRDRPQP
		•		1		WFAWPPRPTRFCGGCWTLTPK
	L	—	L	<u> </u>	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23202	53570	Α	23336	853	1602	PGAAWSRPDLRGCCTEPQPALR
	1					MLVLASPCPQPLAFSSVETMEG
						PPRRTCRSPEPGPSSSIGSPQASS
						PPRPNHYLLIDTQGVP\YTVAW
						LDEESQREPGASGAPGQKKCYS
						CPVCSRVFEYMSYLQRH\RFSH
						SEVKPFE\CDICGK\AFKRSQHL/
						SARHHSIHLGGWVGGPTACRFC
						PRRF\RDAGELAQHSRVHSGER
						PVSVSTLPSPLYGAEHTAETHA
						VEASHEAGLPGAPGTHRTLQG
						ALDSCPDTW
23203	53571	Α	23337	1	1144	MGHHLMIDKLVALGGLYYAIQ
		1				RHYATKCSVLKNDQILVIGLFM
		1			,	IQNVIYRKHFANPLSALFLQQGI
		1				ELFAAIAEIHITTVAERNHAITQI
		1				RLEAQTIFDSLKEWHNAIRKSL
		1				DYEALQAFEFRVGATDRGFPAL
İ						SSEALVRVLVLDANDNSPFVLY
						PLQNGSAPCTELVPRAAEPGYL
		1				VTKVVAVDGDSGQNAWLSYQ
						LLKATEPGLFGVWAHNGEVRT
		1				ARLLSERDVAKHRLVVLVKDN
		1				GEPPRSATATLQVLLVDGFSQP
						YLPLPRAAPAQAQADSLTVYL
			1			VVALASVSSLFLFSVLLFVAVR\
						LCRRSRAAPVGRCSVPECPFPG
						HLVDVSGTGTLSQSYQYEVCLT
22204	52.570	<u> </u>	22220	1,	1260	GGSGANEFKFLKPVIPNLLSRDS
23204	53572	A	23338		1260	MVCYIQENLPFLLKPSVENFYIL
]	ITEGALDREIRAEYNITITVTDL GTPRLKTEHNITVLVSDVNDNA
						PAFTQTSYTLFVRENNSPALHIG
						SVSATDRDSGTNAQVTYSLLPP
		1				QDPHLPLASLVSINADNGHLFA
					İ	LRSLDYEALQAFEFRVGATDRG
l						SPALSREALVRVLVLDANDNL\
						PFVLYPLQNGSAPCTELVPRAA
						EPGYLVTKVVAVDGDSGQNA
		1				WLSYQLLKATEPGLFGVWAHN
		1				GEVRTARLLSERDAAKHRLVV
1						LVKDNGEPPRSATATLHVLLVD
		1				GFSQPYLPLPEAAPAQAQAEAD
				1		LLTVYLVVALASVSSLFLLSVL
						LFVAVRLC\RRSRAASVGRCSM
					1	PEGPFPGRLVD\VSGTGTLFQSY
						QYEVCLTGGSETGEFKFLKPITP
		1				HLPPHRGGKEIEENSTLPNSFGF
<u> </u>	L		<u> </u>	L	<u> </u>	TIET THOUSELEENS TEFNSFOR

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23205	53573	A	23339	123	5794	SSSDSGSLKGACTEVKMEPAGE RFPEQRQVLILLLLEVTLAGW EPRRYSVMEETERGSFVANLAN DLGLGVGELAERGARVVSEDN EQGLQLDLQTGQLILNEKLDRE KLCGPTEPCIMHFQVLLKKPLE VFRAELLVTDINDHSPEFPERE MTLKIPETSSLGTVFPLKKARD LDVGSNNVQNYNISPNSHFHVS TRTRGDGRKYPELVLDTELDRE EQAELRLTLTAVDGGSPPRSGT VQILILVLDANDNA
23206	53574	С	23340	20	313	
23207	53575	A	23341	119	247	RSRCISASPRRPASFRKSIPEEED ESLNI*KGQVSILFFYIK
23208	53576	В	23342	55	288	
23209	53577	Α	23343	1	3576	MAFLFEFSDLVSLLSIGTMLAY SLVAFSVLVLRYQPDENFSKNE KPKEEVVEMNSVPKAESPACVP EASSTPASLWSPVSTIPTPRWGR IVYGCAFLLVVLLSMLCLVLAH WPKRLFSGELIYIAAAVLLLVLI VGFTFTVWRQPQSNTPLYFKPP HISTSQNLSLPEHPMEYLLRTGI EPDVRGVLTVNALPAGPPVACS PLVSISAHVYLMMQMTTETWA QCGVWMLIALECPNLKLKKPP WLHMPLAMTMYA

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	""	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	•	
23210	53578	Α	23344	3	1762	SVRPGPGHVRV\RFFSLSPLPLR
			1	:		PWPLCPAARRTMGSILSRRIAG
						VEDIDIQANSAYRYPPKSGNYF
						ASHFFMGGEKFDTPHPEGHLFG
						ENMDLNFLGSRPSQFPYVTPAP
	1					HEPVKTLRSLVNIRKDSLRLVR
ĺ						YKDDADSPTEDGDKPRVLYSL
1						EFTFDADARVAITIYCQASEEFL
l						NGRAVYSPKSPSLQSETVHYKR
	ļ					GVSQQVSLPSFKIDFSEWKDDE
			1	1		LNFDLDRGVFPVVIQAVVDEG/
		İ				DSVVEVTGHAHVLLAGFEKHM
						DGSFSVKPLKQKQIVDRVSYLL
						QEIYGIENKNNQETKPSDDENS
						DNSNECVVCLSDLRDTLILPCR
						HLCLCTSCADTLRYQANNCPIC
						RLPFRALLQIRAVRKKPGALSP
						VSFSPVLAQSLEHDEHSCPFKK
		ł	ļ			SKPHPASLASKKPKRETNSDSV
		i		f		PPGYEPISLLEALNGLRAVSPAI
						PSAPLYEEITYSGISDGLSQASC
					i	PLAAIDHILDSSRQKGRPQSKAP
						DSTLRSPSSPIHEEDEEKLSEDV
	}					DAPPPLGGAELALRESSSPESFI
						TEEVDESSSPQQGTRAASIENVL
		<u> </u>				QDSSPEHCGRGPPADIYLPALG
23211	53579	Α	23345	35	3134	RPREGPTWPQLPGSSPRRLFPW
						SPGSSSRGLFPWSPGSSSRGLFP
			1			WSPGSSSRGLFPWSPGSSSRGLF
						PWSPGSSSRGLFPWSPGSSSRGL
						FPWSPGSSSRGLFPWSPGGSSR
						GSSRGHPGPPPGGSSRGHPGPPP
						GALPVVTRVLLPGLFPWSPGSS
						SRGLFPWSPGSSSRGLFPWSPGS
						SSRGLFPWSPGSSSRGSSRGHPG
						PPPGALPVVTRVLLPGLFPWSP
						GSSSR/RTLPVVTRLLLSGTLPV VTRVLL
23212	53580	A	23346	3	91	VIKVLL
23212	53581	A	23347	3	174	
23213	53582	A	23348	3	360	
23215	53583	В	23349	158	368	
23213		Г.		1,20	1500	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ļ		sequence		
23216	53584	Α	23350	3786	4928	PPPARTRPCASLTPEKAKWWRS
						KPGLTRAPARCGLSSPQTGSCS
						APASAG*VSGNSRSGTRRGLRP
				1		TRG*GPCGPSSRARAISSPRAS/L
ŀ					}	RMSQRELGLWDPNNFEEPVAL
						QEMDTSNGVLLPFYDPDSSIVY
						LCGKVLTAGQGEQGTGWRCG
						GPCPGAPLNRLILQGDSSIRYFEI
						TDEPPFVHYLNTFSSKEPQRGM
						GFMPKRGLDVSKCEIARFYKLH
						ERKCEPIIMTVPRKSDLFQDDL
						YPDTPGPEPALEADEWLSGQD
						AEPVLISLRDGYVPPKHRELRV
						TKRNILDVRPPSGPRRSQSASDA
1						PLSVRSALLHSGPIYISPNRPYTT
ľ						RPLCPPPQQQHTLETLLEEIKAL
						RERVQAQEQRITALENMLCELV
23217	53585	Α	23351	1	1898	MSRKGPRAEVCADCSAPVGPD
						FALAISELVSPDIFMQSHSENAIS
		ŀ				VKEIVTEIESISQGVGQIQLKGDI
1						LPNPCHTPKKNSIHELLLERAQT
						PENKPGHMEQDEDSCTAQPEL
						AKDSGMCNPEGCLTTHSSIADL
		1				EEGEQLRGNKSSRAQGCTQNS
		1				KNDSSVADLAPKGKSDEAPPEH
		1				SFVLKEPEMSKGKGKYSGSEA
		ł				GSLSHSEQNATVPAPRVLEFDH
		ŀ			}	LPDPQEGPGPEASNGLKVSQKS
1						EGSQPSSSRSPLEFLKEAGVPNQ
1						TLSSGTPDPYFMALLGPKLNSR
						RPHLHPVSHTLDLGNSPPLPQQ
						FAKCQRGRKGPEGKYPLHYLV
						WHNRHRELEKEVRAGQTLITQ
						TLSQANPTAITAEEYFNPNFELG
						NRDMGRPMELTTKTQKFKAKL
						WLCEEHPLSLCEQVAPIIDLMA
						VSNALFAKLRDFITLQIPIFHILN
		1				ARITFGNLNGCDEPVPSVRGSPS
						SETPSPGSDSSSVSSSSSTSEAPR
					!	ENACPSALPGVASCRGCEISPAL
						FEAPRGYSMMGGQREAATRDD
						DDDLLQFAIQQSLLEAGSEYDQ
						VTIWEALTNSKPGTHPMSYE/R
						SPTGQGQVPAATCSG/RYDEQL
						RLAMELSAQEQEERRRRARQE
22219	52504	Ι	22252	341	711	EEELERILRLSLTEQ
23218	53586	Α	23352	341	1/11	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
02210	150.507	<u> </u>	00050	150	710	TOOL MA ENGENIERRI RELEGI
23219	53587	Α	23353	57	719	TSDLMAEIIQERIEDRLPELEQL
		1				ERIGLESHAEIKAIIKKASDLEY
		l				KIQRRTLFKEDFINYVQRIGYSF
						KKDEIENSIVHRVQGVFQRASA
		1				KWKDDVQLWLSYVAFCKKWA
		1				TKTRLSKVFSAMLAIHSNKPAL
		l			:	WIMAAKWEMEDRLSSESARQL
		1			1 1	FL\RALRFHPECPKLYKEYFRM
1		1				ELMHAGKTGGRRRKNLKKPV
						WMWRILIILKKSLRASWHGSST
23220	53588	Α	23354	27	352	TESVVTIMSLLNKPKSEMTPEE
		l				REEEEFNTGPLSVLTQSVKNNT
		l				QVLINCRNNKKLLGRVKAFDR
		l				HCNMVLE\NVKEMLTEG/DRYI
						SKMFL\AGDPVIVVLRNPLIAGK
23221	53589	Α	23355	32	575	STRLDLPKCWDYRSLFNKP\KS
						EMTP\EELQKREE\EEFNT\GPTP
		1				LCLTQS\VKNNTQSAQSTC\RNN
		1				KKLPGAA*RPFDRASPPCSPPGT
		l				CNMVL\ENVK\EMWT*GTQEV
						GKGKEGSSKP\VNKDRLTSSKM
		ŀ				FPGGGGFSSFVVPCGNPFHSAG
		1				QIRGPPVLFDKNSHPLVLLKTA
						CHLVFENY
23222	53590	Α	23356	103	713	VTSGCGKKKVWLDPNET\NEIA
						NANFRSADPGRLIKDGLIIRKPV
						T\VHSRARLPEKTPLARCKGHG
						TLGIGKRKGTANA\RMPKEGHH
						GMRRMRILAPACSRRYRESKKI
		l				D\RHMYHSLYLKVKG\NVFKNK
						ADFSWEHIHKAEGRARPRKEAS
						WLDQAEA\RRSKDQGKHGKRR
						EERLPGQRKEEINQRLYSKEGR
						RPRNKNLTL
23223	53591	A	23357	1	2847	
23224	53592	Α	23358	1	2823	
23225	53593	Α	23359	3	2858	AMFGFQRRGLGTPRLQLWLLL
		1				LEFWEVGSGQLHYSVSEEAKH
İ		l				GTFVGRIAQDLGLELAELVQRL
1		1				FRVASKTHGDLLEVNLQNGILF
1						VNSRIDREELCGQSAECSIHLEV
						IVDRPLQVFHVNVEVKDINDNP
]				1		PVFSLREQKLLIAESKQSDSRFP
						LEGASDADIEENALLTYRLSKN
]					EYFSLDSPTNGKQIKRLSLILKK
						SLDREKTPELNLLLTATDGGKP
						ELTGTVRLLVQVLDVNDNDPE
1				1		FDKSEYKVSLMEN
L		L	i	<u> </u>	<u> </u>	I DIOUTIX VOLIVIEN

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23226	53594	A	23360	194	931	NSPVVPVMASGQFVNKLQEEVI CPICLDILQKPVTIDCGHNFCPQ CITQIGETSCGFFKCPLCKTSVR\ RDAIRFNSAVAEIWWEKIQALQ ASEVQSKRK\EAT\CPRHQEMF HYFCEDDGKFLCFVCRESKDH KSHNVSLIEEAAQNYQGQIQEQ IQVLQQKEKETVQVKAQGVHR VDVFTDQVEHEKQRILTEFELL HQVLEEEKNFLLSRIYWLGHEG
23227	53595	A	23361	2	381	TEAGKHYVEIPLMPTVERSQEA AINWTSARPVSQCWDP\CPPSQ VVCLAGLPP*RVCQNPVKGP\N CPLAHTSEPTGHCLL*TRGMCF PSLCGQLCP*SGPPDCPSPRALH HTGFCPVALLVAESQVRARAC GPEPHRAGRASRWCLHAG
23228	53596	A	23362	213	574	PQTHTGNHSERGRQPITRSPEKP EP\CPSKRAPQTQQP\PWAPSLC PPALGPCKGTRRPRVQVHRPTS SQKCPLPAWFHHLPLWPTSAAR SSSTETPTGQQMRKLRHPAAPR SANALSLGE
23229	53597	Α	23363	21	456	
23230	53598	Α	23364	2	98	
23231	53599	A	23365	64	241	PWPLASNPRGCLP*PSTLN*PTP SSQPGMSLDSLDRHPP*TRETAS GIGGQRPGSDPARTPHFISQTQ
23232	53600	Α	23366	209	353	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23233	53601	A	23367	1766	3292	SSCPTTRVLIMKGAGHPCYLDK
						PEEWHTGLLDFLQGLQLWLMH
						MQQLMSLAVVSGRLSRGQCGT
						ERPSRRAAAAAAAGQPQRGHF
						QHIEERLETNVQNCQSPDTLCP
						THCSPVLNEPCPEVSVVAVAPA
						SPDRMSGSDGGLEEEPELSITLT
						LRMLMHGKEVGSIIGKKGETV
						KRIREQSSARITISEGSCPERITTI
1						TGSTAAVFHAVSMIAFKLDEDL
						CAAPANGGNVSRPPVTLRLVIP
						ASQCGSLIGKAGTKIKEIRETTG
						AQVQVAGDLLPNSTERAVTVS
						GVPDAIILCVRQICAVILEGFSV
						QGQYGAVTPAEVTKLQQLSSH
						AVPFATPSVVPGLDPGTQTSSQ
						EFLVPNDVSRGWDGWRLIGCVI
						GRQG\TKINEIRQMSGAHIKIGN
						Q\AEGAGERHV\TITGSPVSIALA
						QYLITACLETAKSTSGGTPSSAP
						ADLPAPFSPPLTALPTASPLAC
						WAHPMPSPSPTLIGFKAHALLG
						FTTCFPRAAAGLGGLHRQDGSS
23234	53602	В	23368	1	363	
23235	53603	A	23369	486	947	LDGAGLLAPVKAHLQKKGPRL
						GTRGCAMPRLGSTPFLKGQCL
						ANSRNTLEAVSLTESSASRVPP
						APVLGSRRPHSPSPWP/P/YYTP
						VPPPPRMALVPPNMHGQGTVG
						LTSVPLACTVAMLGSYKQAGA
,						FGPLANCYGGLGHAQCWGQEF
22221	52.50	<u> </u>	2225	20	600	СРТОСР
23236	53604	A	23370	38	680	GSRLRRLQAAAARPALPLPLPP
						WEWKHLPHVPEAKWWLTTAR
						HSAAYRADPLRVSSRDKLTEM
						AASSQGNFEGNFESLDLAEFAK
						KQPWWRKLFGQESGPSAEKYS
						VATQLFIGGVTGWCTGFIFQKV
]		GKLAATAVGGGFFLLQLANHT
						GYIKVDWQRVEKDMKKAKEQ
						LKIRKSNQIPTEVRSKAEEVVSF
22227	52605		22271	20	200	VKKNVLVT\GGFFGGFLLGMAS
23237	53605	C	23371	30 14	200 161	
23238	53606	Α	23372	14	101	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23239	53607	А	23373	69	715	APALPGCEHMMAIRELKVCLL
						GDTGVGKSSIVWRFVQDHFDH
		1				NISPTIG\ASFMTKTVP\WENDFT
		1				KFLIWDTAGQERFSFIGLPLYYS
ľ		1				RLQLAAVYSCYGYLPSRDFIFIT
		l				LKE\WV\KRLKELGPE\NIVMAI
						AGNKCDLSDI\REVPPEGMLKE
						Y\AESIGAIVVETSAKNAINI\EE
ŀ		1				LFQGIS\RQIPPLDPPMKNGKQW
ļ						DHQKLRSQPMQASRRCC
23240	53608	Α	23374	3	293	VTSTMAYRAQGQKV\QKVMV
ŀ	i					QPINLIFRYLQNRSRIQVWLYEQ
						VNMRIEGCIIGFDEYMNLVLDD
		l			:	AEEIHSKTKSRKQLGRIMLKGD
		İ				NITLLQSVSN
23241	53609	Α	23375	3	393	HESCSQRQRAGCALCENSHPW
						RSRGPGVQKSAEGLWCQPICRT
						SSSRYL\QNRIAGFQVWLYEQV
						NM\RI\EGCIIGF\DEYVNLVLDD
		İ				AEEIHSKTKSRK\QLGR\IMLK\G
					:	DNITLLQTCLPPRMVHGSGEIV
23242	53610	С	23376	129	425	
23243	53611	Α	23377	470	635	ENSSSHHIIKPFSSGCSIHGFSGH
		l				INVTSVTIL*PIKSCHIRCHF*CA
						QYMVADN
23244	53612	Α	23378	71	566	PLFPPCLSPVVAGWARGHMLG
						QSHQQVAAPLKHRPITPTELCQ
		l		1		KCLSLIKKTKFLNAYITVSEEVA
						LKHAEESEKRYKNGQSLGDLD
						GIPIAEKDNFSTSGIETT\CASNM
		ł				LKGYIPPYNATVDQKLLDQGA
						LLMGKTKLNDFAMGSGSTDGV
		<u> </u>				FGPVKNPWSYSK
23245	53613	Α	23379	255	1042	ILAARMGKQNSKLRPEVMQDL
1						LESTDFTEHEIQEWYKGFLRDC
		ļ				PSGHLSMEEFKKIYGNFFPYG\D
						ASKFAEH\VF\RTFDANGDG\TID
1						FREFIIALSVTSRGKLEQKLKW
1		1				AFSMYDLDGNGYISKAEMLEIV
		l				QAIYKMVSSVMKMPEDESTPE
						KRTEKIFRQMDTNRDGTWVPH
						LALIYRTAEGGFAVQRASVSFY
]		PPPSAGNAMVNEAAVFLEIPLG
			1			SSPGQVSAVCHSGRKLSLEEFIR
						GAKSDPSIVRLLQCDPSSAGQF
	l		L	L	L	7

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23246	53614	l A	23380	1	721	 MDGNLRKIRFEKHAITEEFRTT
23240	33014		25500	'	/21	EPGLAALRLPSTWIRGVRTSES
						KREVGKGGQSQQYENRDTEGE
				:		GHVMTKARDWSKAAASQRLP
Ì					İ	ANHLKLRLCIDGFRSHSLPEEGP
1						KKPCQHHAAQEEPCTRSGRLPQ
1						AHEGSISDCCTSSEQGSMDIGSV
l						EPGTGYNLLVCHLL/STLEKRSI
i					ļ	WAGVSRFSRYHLSRLPLARK\G
E						NPLTPCASQLLLPLPSTTQKRQS
ŀ						GHALYSFPYRPQPKNDNLVMP
23247	53615	A	23381	80	562	DRAMWGDLWLLPPASANPGT
202.7		1	23301			GTEAEFEKAAEEVRHLKT/RSPS
						DEEMLFIYGHYKQATVGDINTE
						RPGMLDF\TGKA\KWD\AWNEL
i						KGTSKGDAVEA\YLNQV\EELK
			ļ ,			KKYGDMERLDLVTVPCVYPKP
						ETIPFFFSNTVDGWEFRENNPV
						KPATQRLLTHTGS
23248	53616	Α	23382	380	1041	EVTINSGFIFIMLFMEHPEHYTI
						MHKWILTWILPTLLYRSCFHIIC
						LVGTISLACND\MTPEQMATNV
i		1				NCSSPERHTRSYDYMEGGDIRV
ļ						RRLFCRTQWYLRIDKRGKVKG
		1				TQEMKNNYNIMEIRTVAVGIVA
						IKGVESEF\YLAMNKEGKL\YA
						KKECN\ED\CNFKELILENHYNT
		l				YASAKL/WTHNGGEMFVALNQ
						KGIPVRGKKTKKEQKTAHFLP
						MAIT
23249	53617	Α	23383	127	549	SSPKMAATMKKAAAEDVNVTF
						EDQ\QKINKFARNTSRITELKEEI
						EVKK\KQLQNL\EDACDDIMLA
						DDD\CL\MIPY\QLGDVFISHSQE
						ETQEMLEEAKKNLQEEIDALES
						RVESIQRVLADLKVQLYAKF\G
						SNINLEADES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23250	53618	A	23384	1	2050	MEYESMVISFDDRVMEKTEYG
						PIFTVFAMGNRMTFVTEEEGIN
	[VFLKSKKVDFELAVQNIVYRTA
						SIPKNVFLALHEKLYIMLKGKM
						GTVNLHQFTGQLTEELHEQLEN
						LGTHGTMDLNNLVRHLLYPVT
						VNMLFNKSLFSTNKKKIKEFHQ
						YFQVYDEDFEYGSQLPECLLRN
			}			WSKSKKWFLELFEKNIPDIKAC
						KSAKDNSMDSDEDPVNLQSRS
						ALVCEMLHVQRSSSDADAIGPH
						ADHSLSNRWEVILADEHGIIAD
						DMISRYALSEKSQVELSTQVIKS
ļ						ASSKSVNKSDIDTSVFLNWYNP
ł		1				EKKMIRAYATVIDGPEYFWCQF
ļ		-				ADTEKLQCLEVEVQTAGEQVA
	1					DRRNCIPCPYIGDPCIVRYREDG
		1				HYYRALITNICEDYLVSVRLVD
						FGNIEDCVDPKALWAIPSELLS
ŀ		ł				VPMQAFPCCLSGFNISEGLCSQ
						EGNDYFYEIITEDVLEITILEIRR
						DVCDIPLAIVDLKSKGKSINEK
		ŀ				MEKYSKTGIKSALPYENIDSEIK
						QTLGSYNLDVGLKKLSNKAVQ
1						NKIYMEQQTDELAEITEKDVNII
						GTKPSNFRDPKTDNICEGFENP
						CKDKIDTEELEGVSQKAQESMC
İ		ŀ				TEDMRKSSCVESFDDQRRMSL
						HLHGADCDPKTQNEMNICEEEF
						VEYKNRDAISALMP\FSLRKKA
						VMEASTIMVYQIISQLNYRTPTL
23251	53619		23385	3	602	TRPPTRPGLSPDPTSPTEADRGV
23231	33019	A	23363	3	002	CERPHRCFLLPQITHPSSGRRCL
						CLLSPTSFDFSLLYESQSDRAIF
						GVGTEPDQEPPPPYHEQVPVSV
						YHPT\P\SQTWVATQLTEEEQIRI
						AQRIGLIQHLPKGVY\DPWRDG
						SEKKIRECVICMMDFVYGDPIR
		1	,			FLPCVHEWYHLDCIDDWLMRS
		İ		·		i I
22252	52620	 	22206	22	510	FTCPSCMEPVDAALLSSYETN
23252	53620	Α	23386	33	512	TAARCQGGPTHVPAFVGTACP
						TRIPGHAPARQCREYYY*YYYD
						F/WKEPRRCHALLLPPWAGLEV
						GMPPSHTQLGMHPAPPPPHLEV
						KAQG\PSNRQRTDGCRWCSGR
,						ASALGGGRQDSRLRERGGGYD
						H/HCPICRSLKIQLKCHPASGPQ
		<u> </u>				TVGLGRLASVS
23253	53621	A	23387	1	1731	
23254	53622	В	23388	1	706	

sequence 09/540,217 co	cation of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
sec	don for nontide		-stop couon, /-possible nucleonue
	• •	of peptide sequence	deletion, \=possible nucleotide insertion)
	quence		
23255 53623 A 23389 I		1986	MGIGEGPTTSEGCRESVEYSHQ
			KSELPLSGDSQVPGPIGCNCDP
			QGSVSSQCDAAGQCQCKAQVE
			GLTCSHCRPHHFHLSASNPDGC
			LPCFCMGITQQCASSAYTRHLIS
			THFAP/GDFQ/GFA/LVNPQRNS
			RLTGEFTVEPVPEGAQLSFGNF
			AQLGHESFYWQLPETYQGDKG
			EAYFARMRRAHQNRTSLSEEQ
			LRAAVTAGRIPEPPEGRDWAQR
			ASQFSLSYEGFSLLPGSLYYWQ
			LPRAFLGDKVAAYGGKLRYTL
			SYTAGPQGSPLSDPDVQITGNNI
			MLVASQPALQGPERRSYEIMFR
			EEFWRRPDGQPATREHLLMAL
			ADLDELLIRATFSSVPLAASISA
	ļ		VSLEVAQPGPSNRPRALEVEEC
			RCPPGYIGLSCQDCAPGYTRTG
			SGLYLGHCELCECNGHSDLCHP
			ETGACSQCQHNAAGEFCELCA
			PGYYGDATAGTPEDCQPCACP
			LTNPENMFSRTCESLGAGGYRC
			TACEPGYTGQYCEQCGPGYVG
			NPSVQGGQCLPESSELHFPSVQ
			PSDAGVYICTCRNLHQSNTSRA
			ELLVTEAPSKPITVTVEEQRSQS
			VRPGADVTFICTAKSKSPAYTL
			VWTRLHNGKLPTRAMDFNGIL
			TIRNVQLSDAGTYVCTGSNMF
	•		AMDQGTATLHVQGSAEAGERS
			MYSISVLLKYTS
23256 53624 A 23390 3		13319	PASGRLRAARSGRRGASERESG
			AGRAMGWRAPGALLLALLLH
			GRLLAVTHGLRAYDGLSLPEDI
			ETVTASQMRWTHSYLSDDEYM
			LADSISGDDLGSGDLGSGDFQM
			VYFRALVNFTRSIEYSPQLEDA
			GSREFREVSEAVVDTLESEYLKI
			PGDQVVSVVFIKELDGWVFVE
			LDVGSEGNADGAQIQEMLLRVI
			SSGSVASYVTSPQGFQFRRLGT
			VPQFPRACTEAEFACHSYNECV
			ALEYRCDRRPDCRDMS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23257	53625	A	23391	41	1323	HYLVFFLPSLRRYTVSLSPNSTS YQLLVPGPSY/SALPISSAASLY AGA\GGSGSRMSVSHSTSLRDG ISSGGLA/TGMARGLGGMGGIQ NKKETTQSLNDSLAVRVRSLET ENRKLESKIREHLEKKGPQVRD WSHYFKTIEDLRAQIFANTVDN VRIFLQIDNARLAADDFRVKYE TELAMRQSVESDIYGLHKVIDD TNVINGCSFFMKKNHGEEVKC LQAQIASSGLTMEVDAPKSQDL AKITADISTRYDELARKNREEL DKYWSQPIEESTTEVTMQSAKV RAAEMTFMELRRTVQSLEIDLD LMRNPKASLENSLREVEACYA LQMEQLNLDP/GILLHLESELTQ TRAEGQL\QAQKYKALLNIKNK LEAEIATYCLLLEDGKDFNLGD TLDSSNPMQTIQKTTTSRIVDG
			2222			KVVSETNDTKVWRH
23258 23259	53626	A A	23392 23393	732	771 1656	SCSWTSPAAWPPNASLHSCLKR
						RNQDHLLLWDWRFLDFPSSMA SKCQSPLLPQKKKPRPLPALGP EKTSASAGLPKKGEKEQQETIE HIHEVQNEMDRLNEQAK\RSEL ISKIPDFWVTIFVNHPEVSALPG EEDEEALHYLTRVEMTDFDDIK SGYRIDFYFDENPYFENNILAKE FHLNESGDPSSKSTEIKWKYGK DLMKRSSQTQNKAHRKRQHEE PESFFTWFTDHSDVGADELGEV IKDIWPNPLQYYLVPDMDNEEG EGEEDDDDDEEEEGLEDIDEEG DEDEGEEDEDEGEKGEEDEGE
23260	53628	A	23394	3	923	ARRLVVWFWDFPNSMAPKRQS PLPPQKKKPRPPPALGPEETSAS AGLPKKGEKEQQEAIEHIDEVQ NEIDRLNEQASEEILKVEQKYN KLRQPFFQKRSK\LSAGIPNFWV TTFVN\HPQVS\ALLGEEDEEAL\ HYL\TRVEVTEFEDIKSGYRIDF YFDENPYFENKVLSKEFHLNE\S GDPSSKSTEIKWKSGKDLTKRS SQTQNKASRKRQHEEPESFFTW FTDHSDAGADELGEVIKDDIWP NPLQYYLVPDMDDEEGEGED DDDDEEEEGLEDIDEEGDEDG EEDEDDDEGEEGEEDEGEDD

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
23261	53629	A	23395	732	1656	SCSWTSPAAWPPNASLHSCLKR
		1				RNQDHLLLWDWRFLDFPSSMA
						SKCQSPLLPQKKKPRPLPALGP
						EKTSASAGLPKKGEKEQQETIE
]		HIHEVQNEMDRLNEQAK\RSEL
1						ISKIPDFWVTIFVNHPEVSALPG
						EEDEEALHYLTRVEMTDFDDIK
	Ì					SGYRIDFYFDENPYFENNILAKE
		1				FHLNESGDPSSKSTEIKWKYGK
1						DLMKRSSQTQNKAHRKRQHEE
	1					PESFFTWFTDHSDVGADELGEV
ļ						IKDIWPNPLQYYLVPDMDNEEG
						EGEEDDDDDEEEEGLEDIDEEG
						DEDEGEEDEDEGEKGEEDEGE
23262	53630	A	23396	3	924	ARRLVVWFWDFPNSMAPKRQS
23202	33030	` `	23370		, ,	PLPPQKKKPRPPPALGPEETSAS
		1				AGLPKKGEKEQQ\EAIEHIDEVQ
						NEIDRLNEQASEEILKVEQKYN
		1				KL\RQPFFQKRSK\LSAGIPNFW
						VTTFVN\HPQVS\ALLGEEDEEA
		İ				L\HYL\TRVEVTEFEDIKSGYRID
ļ						FYFDENPYFENKVLSKEFHLNE\
1				}		SGDPSSKSTEIKWKSGKDLTKR
						SSQTQNKASRKRQHEEPESFFT
						WFTDHSDAGADELGEVIKDDI
						WPNPLQYYLVPDMDDEEGEGE
	1			1		EDDDDDEEEEGLEDIDEEGDED
				İ		EGEEDEDDDEGEEGEEDEGED
23263	53631	A	23397	1	1393	MGKQKGGGPQAWEAEGLGRE
23203	12021	A	23371	1	1393	WMRMPQEGKGSQRKSTQKQT
						KREIDAYIVQAKERSYETVLSF
						GKRGLNIAASPAVQAATKVLW
						APSPPAAPIPPLRPFGLIQLLCQG
						AQRWQPGSREVKGRLCQARLA
						ETSSIAGNLRVQSYERPVDPGV
						TMDPLLPGRTIVFAAGDHTLW
						RAPVSSCAQRSGVWSLDNMGV
						NSGSVPFHGVRIQASQLISLSFN
			ļ	1		FSFCEADLKTPLHGFLILPAGPK
						EEVTVKKEKRERDRDRQREGH
						GRGRGRPEVIQSHSIFEQG\PAE
		İ		1		MMKKKGNWDKTVDVSDMGPS
						HIINIKKEKRETDEETKQILRML EKDDVDHHALSEHLVTTATFSI
:						GSTGLVVYDYQQVECGGHVRA
	1					DLKTQDLYSHAQFGDNNYPGG
				1		VDWYRAGGLQDSDTEDECWS
		1		1		DTEAVPRAPARPREKPLIRSQSL
				1		RVVKRKPPVREGTSRSLKVRTR
2000	50.522	4.	22222	1007	504	KKTVPSDVDS
23264	53632	A	23398	297	584	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23265	53633	IA	23399	2	149	CMKFGAVTPIGGPPLGDQSPVL
1000		` `				LLFAL*KRSTYNLGSSDPPAQGI
		1				SHQF
23266	53634	A	23400	1	170	PAGIRHEERSTYDLWSSDRPGQ
				<u> </u> ^		GTSHQF*IIRFQIPWNSGPRLSE*
						LLPRSSRLSS
23267	53635	A	23401	1	1077	
23268	53636	В	23402	289	1359	
23269	53637	A	23403	3	2233	
23270	53638	A	23404	1	1491	
23271	53639	В	23405	76	511	
23272	53640	Α	23406	1	2706	
23273	53641	Α	23407	273	862	
23274	53642	Α	23408	1	1080	
23275	53643	Α	23409	134	675	MHYFHSSVQIGLPLTRRSQVLIL
						PQPLLPNPIILLSPPLLTPVPAYS
ŀ						FIL/PTSPPPPARQFPLKKVAEAK
						GIVKQLKTDTARSPRKPTGPSQ
				:		TLWVTLTVEDLKHAFFTIPLHP
						SSQPLFAFTWTDPDTHQAQQIT
				ł		RAVLPQVFTDSPHYFSQAQISSS
						SVTYLGIILIKTHVLSLLIMSG
23276	53644	Α	23410	1	895	MEVHKHPFNTLALRHESGCEL
				}		VWAEKHRNQIFPFPKSTASPKQ
						WDTGRHNSDAKQQEPGGREER
			ŀ			NTMEMETKETHFIRGPKTLASV
						TDWEGSLPLVFNHCRGTSLIIHP
						RFKGVRPRRDACLGPSPLAASP
						AFLGKGQHALKGLKPVITRLLQ
						HGLLKPINSPFPS\PILPVLKPDK
						AYKLVQNLRLINQIVLPIHPVVP
1						NPYTLLSSIPPSTTHYSVLDLKH
						AFFTIPLHPSSQPLFAFT*TDPDT
			İ			H*AQQITWAVLPQGFTRQPPLL
						QLSPNFILICYLSRHNSHKNTRA
						LPADRV
23277	53645	Α	23411	1	930	MIQEHADKQVQRLQGVLLGSIP
						TAASRARLHPGEINNHVAHTEP
				;		VWWSLHMDAYEIWCRGSDRR
						TSLGRSIPLPPALCSLRRIHLRPQ
						VLRSTSPRNISPISNPGHLSDYTP
						TFQGCQTMQGRLPWSFTLSGK
						SRFSGEGATQRQYPIPQQALKG
						LMPAITRLLQHGLLKPINSPYHS
		1	1			PILPVLKPDKPYKLVQDLRLINQ
		1				IVLPIHPMVPNPYTLLSSIPASTT
						HYSV\LYVKHAFFTIPLHPSSQP
						LSTFTWTDPDTHQAQQIT*AVL
						PQVFTDSPHYFSQAQISSSSVTY
1	I	1		1		LGIILHENTRALPADHV

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	I .	•	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23278	53646	A	23412	299	1028	RLGSFSSNHPYLGQRQYPIPQQ
						ALKGLKPVITCLLQHVLLKPVN
				-	İ	SPYNSPILPVQKLDMPYRLVQD
						LCLINQIVLPIHAMVPNPYTLLT
						SIPPSTTHYSVLDLKPAFFTIPLR
	1					LSFQPLFTFSWTDPDTHQAQQI
		İ			İ	TRAVLPQSFTDSPHYFSQAQIS*
	•					RLGSFSSNHPYLGQRQYPIPQQ
						ALKGLKPVITCLLQHVLLKPVN
	1				}	SPYNSPILPVQKLDMPYRLVQD
1	1					LCLINQIVLPIHAMVPNPYTLLT
						SIPPSTTHYSVLDLKPAFFTIPLR
	İ					LSFQPLFTFSWTDPDTHQAQQI
						TRAVLPQSFTDSPHYFSQAQISS
						SSVTYLGIPGFAILTKPLYKLTK
ļ						GNLADPIDPKSFPHSSFRSLKTA
		ļ				LETAPILALPVSSQPFSLPTAEV
		<u> </u>				WAAQSEFLHKNRDRTE
23279	53647	Α	23413	157	202	RSECARWMEK*CPWL
23280	53648	Α	23414	184	482	EGVSVFLLLSPYQGQPQVSRRC
						S/RQVSQRCPHRGRGPRCPQVS
						RSHGPSWVGHPGMPLYGS\EQF
						YCGAPGLAGPPNGSRGESLKIS
						VWIFPCRDFGNFN
23281	53649	В	23415	13	862	
23282	53650	Α	23416	1	11342	MATRWFYYGHPPRDARVPRIV
						AEAKEKNLTPGKIIEVDVTIEAQ
					1	IRQEFLDAERDTGGFVKFFRITK
						KKLQEFFLGEEEEQVPHVSPVE
ļ						LGAKPTAFNAFIANAVGKTLAV
ļ						QHASLKAPEPEMTVTHERFYG
						NYAVWITDVDPRNHRIEVIHVI
						REISGRTSEYVNNLLAQPTPFK
						VFDWLTETQAKEFVRRLNNTQ
						GVISVYNKLEDGIPVKVAVKAT
						EPTNKEKLEAASTKLKAMGAV
		<u> </u>				ATDAYEFHTDLNENE
23283	53651	Α	23417	7597	7883	LSSLAWVWQWNHRISVLFHSN
						SVRTNLVPQFNQLVITESNSCPE
ĺ		i				FFNQTSAFFKT*VESGCM/CLQD
						*GIEWFRLRLDVYNSRLVFLTG
						ATDYCTG
23284	53652	Α	23418	3	1573	
23285	53653	A	23419	6692	6793	GASVAPSIGS*STGNAIPSCLFQ QNVIVPIPQS
23286	53654	Α	23420	1	17727	
23287	53655	Α	23421	6687	6785	VNTGNRTWWEPLFRK*KKTQR
						EDTTEKKTKGR
23288	53656	Α	23422	1	2856	
23289	53657	Α	23423	1	7194	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23290	53658	A	23424	7460	7561	GASVAPSIGS*STGNAIPSCLFQ QNVIVPIPQS
23291	53659	Α	23425	1468	1857	
23292	53660	Α	23426	1	1985	MEPVGIDIIAISWIGAPEYPDNV
						KRAINAKVEATQKTLQREQEIQ
		1				QRVAEANMEREQAKGVADAIR
						LRGEALTENPNVMQLEAINKW
						NGLVPFITSSPGMGKSAIVRQIA
		ĺ				EDFGLKLIDHRLSTSAPEDLSGL
						PRFKENGRAEFAPFDELFPLEG
						DAVPEGFNGWLLFLDEFNSAR
						KEIQAAAYKLVLDRMTGQKKL
						HPYVAIVCAGNKATDRAITNNL
						STAMQSRLIHLEMETDFDVFME
		1			1	DVAIPNKWDERVIAFLNANPNK
						LNDFEPDHQEKTFCCSRTWEFV
		ŀ				NKIVSILPPGPINSEMTLLLAGTI
		İ				TSGVATSFVQFTQVYSNMVSL
		1				NEILDNPKKARMPEDNNLLWA
						VVTSLINNTDEDNHSKIFDYVE
						RMPFTFKVLYYRSRKTDMSMT
						DAQLTREYDRCQAKAFMGKTA
						AFFGSLLCSLKFRWVKDGCDT
		1				AQTDGEHLEWNPDWFESLLPE
		1				SRVTVLMHELWHVGLLHSVR
		1				MGSRDPEVWNYACDIYINNQLI
						QDGYSFIGIENCWKDPKYAGW
						AEEQIYHDLMSKHPRPPKASGA
						FGTGSGGDMKPSTSKASLKLSG
	1	1				GSLPGVLPGKMEEVITQFLKPV
						VPWQVLLERFFNDLQETYYSW
	·					QRPNRRYPDMYLPSSMDDDCR LEHLAYFL*NEADLTGLLVLK*
						TDDGQSVSGQNLLQSDEQLQIY
23293	53661	A	23427	2365	2784	TDDOQ3 V3GQNELQ3DEQLQTT
23293	53662	A	23428	5789	5980	DQNMRFDNTFETYYWKPTIIVII
23294	33002		23420	3767	3700	IRKLLMNDSSKDTLK*LRCWK
						NWLQILKKHTIHFIMPLQL
23295	53663	A	23429	1	6819	I W EQIBERRITIII I I I I EQE
23296	53664	A	23430	376	2052	SKVCGNDGFADDLKHLLIFTGA
23270	33004	`	25 150	370		QSGEDIVPFQLEDGHSLGEVVL
		İ				LHGGSGVESGQR/CGRTSAGSC
						Y*SPGKICCCNLLVTKTEGFFAP
			<u> </u>		1	VPIQLENTNQVTNCSDNW
23297	53665	Α	23431	2	2189	
23298	53666	A	23432	2	1583	
23299	53667	A	23433	2142	2242	GASVAPSIGS*STGNAIPSCLFQ
	1				}	QNVIVPIPQS
23300	53668	A	23434	3	1188	
23301	53669	A	23435	470	862	
23302	53670	A	23436	1	1352	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23303	53671	A	23437	1837	1955	SKCSIGLYQRYPYSNGC*NSGCI CRVY*CCREHRGYVRN
23304	53672	A	23438	1447	1565	SKCSIGLYQMYPYSNGC*NSGC ICRVY*CCREHRGYVRN
23305	53673	A	23439	773	823	
23306	53674	A	23440	3	1529	
23307	53675	Α	23441	1	1794	
23308	53676	A	23442	236	427	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23309	53677	Α	23443	8510	8611	GASVAPSIGS*STGNAIPSCLFQ QNVIVPIPQS
23310	53678	A	23444	516	669	ATSELKRKASLGLMEPDVSRK* ARCSSLPSSSIGDGRYISGYLRL GRCHE
23311	53679	A	23445	410	593	FLNLIQPQGKRKLLLKRCRSLLP KN*CQHSKQHKLSISSVVILNIT NS*PNKVKLMPILAM
23312	53680	В	23446	1	2286	
23313	53681	A	23447	1418	1723	CNTNIIFNCHFPTKLGCICKAMK VVSPLIHNSIVILLNQITIDFFSTL AQICTFICLTYSKQTTTRIASFK KVIEEFLSLAPIEKVELKEHS*YI SISEKV
23314	53682	В	23448	1	6167	
23315	53683	Α	23449	1	3199	
23316	53684	Α	23450	1	5733	
23317	53685	Α	23451	709	1099	
23318	53686	A	23452	5039	5230	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23319	53687	A	23453	664	993	
23320	53688	A	23454		19859	MASIDSIAVCNPRQVRKFVEHC IRSGLVPFITSSPGMGKSAIVRQI AEDFGLKLIDHRLSTSAPEDLSG LPRFKENGRAEFAPFDELFPLEG DAVPEGFNGWLLFLDEFNSAR KEIQAAAYKLVLDRMTGQKKL HPYVAIVCAGNKATDRAITNNL STAMQSRLIHLEMETDFDVFME DVAIPNKWDERVIAFLNANPNK LNDFEPDHQEKTFCCSRTWEFV NKIVSILPPGPINSEMTLLLAGTI TSGVATSFV
23321	53689	В	23455	74	240	
23322	53690	Α	23456	143	384	
23323	53691	Α	23457	1	3267	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23324	53692	A	23458	2	550	KRSSEHIQRIDRDVSGTLRKHIF FRDRYGTKQRELLHILLAYEEY NPEVGYCRDLSHIAALFLLYLP EEDAFWALVQLLASERHSLQG FHSPNGGTVQGLQDQQEHVVA TSQPKTMGHQDKKDLCGQCSP LGCLIRILIDGISLGLTLR\LWDV YLVEGEQAFMPITRIAFKVQQK RLTKTSR
23325	53693	Α	23459	72	630	
23326	53694	A	23460		2835	MDALMLALGALMLALGLGDP KSAQVSDPDGAEPGPGRATGT DPLVQIHWKLGISEALSIPGGRL YRDLWSLTPVCLHIPGIAHHGP FTLGRMDVVEVAGSWWAQER EDIIMKYEKGHRAGLPEDKGPK SFGSYNNNVDHLGIVHETELPP LTAREVKQIRREISRKSKWVKM LGEWDTYKNSRKLIDRAYQGIP MNIRGPMWSVLLNIEEIKLKNP GRYQVRSARAQPTGQAVSGAQ VSSWRERQDHPGELGVKI

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SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23327	53695	A	23461		832	MGWRARWLMPVISALWEAED PIPPELTRLLQQSQDPLLMGLFP TNPKEKTQEEPPGQSRAPVLTV VSKFKVLAGLLHLGNIQFAASE DEAQPCQPMDDAKCEGQGVG WAGPVRKASLEQLLQVLHSTT PHYIRCIKPNSQGQAQTFLQEE AGAGRRQRRQKRQKQQRRRRQ LGGGGDGGNGSVGDDGATRGS RGCALEKECVATPRRGLLFSRP HKLAFAVGGSVTSVGPKPRAP MDVVEVAGSWWAQEREDIIM KYEKGHRAGLPEDKGPKPFRS YNNNVDHLGIVQDFADFGTTIK QDFRLLGQTSVDRLLQLSQGQ AVKGNQLLPVSLVKRKTTLAP NTQTASPRALADSLMQLARQV SRLESGHFYDAHNPFLPPPSMS KQAKKTGVPSEGLCRPGGEGQ SAVARLTGEKKNHPGAQYANR LSPRVGRFINAAGTTGFPTGKR AVSATQLIGLQLILLGEVLQKV LTASYPAQSSVPLTSGPVRQIM KEKGKRSSEHIQRIDRDISGTLR KHMFFRDRYGTKQRELLHILLA YEEYNPEVGYCMDLSHIAALFL LYLPEEDAFWALVQLLASERHS LQAARAPAAIGAHERADQAQIS LGLTLRLWDVYLVEGEQALMP ITRIAFKVQQKRLTKTSRCGPW ARFCNRFVDTWARDEDTVLKH
23328	53696	A	23462	3	497	LRASMKKLTRKQGDLPPPAKPE SSECLTMAWIPLLLPLLILCTVS VASYELTQPASVSVSPGQTASIT CSGDVLAKKKYARWFQQKPG
						QAPVLVIYKDSERPSGIPERFSG SSSGTTVTLTISGAQVEDEADY YCYSAADNNGWVFGGG\TKLT\ VLSQPKAAP\SVTLFP\PSSEELQ ANKAHTGVSQ
23329	53697	A	23463	1	393	FRSGSELVKAGLRAFFENAAED LEKTSENLKLGKFTHSRTQIKG VSQNINYTTVALLPILTSIFEHV TQHQFGMDLLLGDVQISCYHIL CSLYSLGTGKNIYVE\ILGMPDT VEDMCPDIPQLECLMKEIN
23330	53698	Α	23464	319	508	
23331	53699	С	23465	94	222	
23332	53700	A	23466	1	636	
23333	53701	A	23467	1	483	
23334	53702	Α	23468	1	1797	

SEQ ID	-		SEQ ID NO:			Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	deletion, \=possible nucleotide insertion)
	Sequence		05/1540,217	sequence	peptide sequence	december, it processes are made insertion,
2222	50500		100460		14076	
23335	53703	A	23469	1	4275	
23336	53704	Α	23470	1	1848	+ DOWN DRIVE TO THE PROPERTY OF
23337	53705	Α	23471	122	14928	ARSWRRRWRTRTKRRRKSAA
		1				AVSARRAAAGGSRGAGGWGT
						ADASGAMAEGGEGGEDEIQFL
						RTEDEVVLQCIATIHKEQRKFC
		İ				LAAEGLGNRLCFLEPTSEAKYIP
		l				PDLCVCNFVLEQSLSVRALQEM
						LANTGENGGEGAAQGGGHRTL
						LYGHAVLLRHSFSGMYLTCLTT
						SRSQTDKLAFDVGLREHATGE
		l				ACWWTIHPASKQRSEGEKVRIG
	1		ļ			DDLILVSVSSERYLHLSVSNGNI
						QVDASFMQTLWNVHPTC
23338	53706	Α	23472	134	1063	RPPGLTRSPSSPAPARSPTRRQS
		1				TRRRSGATTSCSQRLWATWNG
ŀ						CD/VCLNQSLREIPTDDKGFTAI
ļ		ļ				HFAAQWGKLACLQVLVEEYKF
						PVDLLTNNSQTPLHLVIHRDNT
İ						TVALPCIYYLLEKGADLNAQTC
						NGSTPLHLAARDGLLDCVKVL
						VQSGANVHAQDAMGYKPIDFC
1						KIWNHRACARFLKDAMWKKD
						KKDFAREMTKMKMFKSQLTL
		Ì				MEHNYLIEYQKEHKILREAAIR
						KWLHGKLHPGHSLVSNTKQAR
						ATALSKTPEQRESQRSRSFHPSV
				1		DARLQCIPQPTEMPKPIYRKPTV
						KRPHNVEC
23339	53707	A	23473	1	1280	MAGLGFWGHPAGPLLLLLLV
					1	LPPRALPEGPLVFVALVFRHGD
1						RAPLASYPMDPHKEVASTLWP
ļ						RGLGQLTTEGVRQQLELGRFLR
1						SRYEAFLSPEYRREEVYIRSTDF
}		l				DRTLESAQANLAGLFPEAAPGS
	·	l				PEARWRPIPVHTVPVAEDKLLR
		ļ				FPMRSCPRYHELLREATEAAEY
					ļ	QEALEGWTGFLSRLENFTGLSL
						VGEPLRRAWKVLDTLMCQQA
						HGLPLPAWASPDVLRTLAQISA
						LDIGAHVGPPRAAEKAQLTGGI
						LLNAILANFSRVQRLGLPLKMV
						MYSAHDSTLLALQGALGLYDG
1						HTPPYAACLGFEFRKHLGNPAK
						<u>'</u>
		1		1		DGGNVTVSLFYRNDSAHLPLPL
						SLPGCPAP\CPLGRFYQLTAPAR
1						PPAHGVTCHGPYEAAIPPAPVV
1						PLLAGAVAVLVALSLGLGLLA
		<u> </u>	<u> </u>	<u> </u>	1	WRPGCLRALGGPV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23340	53708	A	23474	133	672	AVGWESGKWDSGRGQGGGQE
						QMWSCQSDMLYPPDQVGAWG
		ļ				LPGPTGPKGDAGSRGPMGMRG
						PPGECPQCCPTAPLAIPSTRWAL
		İ				PCSWPPGPPGPPGPPAPVGPPHA
		ŀ				\GSPSMVSPPGIPAGGGGGVAI
						STVPATIKTRRQEVSPELPRP/PG
						PAPPPGPDVRPWLLRPPGPMGK
						LSPGQG
23341	53709	A	23475	135	677	AVGWESGKWDSGWGQGGGQE
		ļ		}		QMWSCQSDMLYPPDQVGAWG
						LPGPTGPKGDAGSRGPMGMRG
		1				PPGECPQCCPTAPLAIPSTRWAL
						PCSGPPGPPGPPAPVGPPHA
						\GSPSMVSPPGIPAGGGGGGVAI
		ŀ				STVPATKKTRRQEVSPELPRP/P
j		İ				GPAPPPGPDVRPWLLGPPGPMG
		j				KLSPGQG
23342	53710	Α	23476	220	745	TVIQRTGLHSVGPGDPQQGLRE
}		ļ				AKLGTMVRLVLLPTPRRTTEVR
						TGPERELSMVLEAILGSLLETAC
						DHRQTPPSSKPQA/GSSRLREGT
						SAGQPCS/LRLRLGPRAGGRRC
						CRDIERSSSSSVCGRKCKRKKR
						KRGMCQSRHKHGCRRPQEGPR
						CNDGRALCSPRPQTTPGPLSTA
23343	53711	Α	23477	1	2433	
23344	53712	Α	23478	3	803	FPGRRFRVRACVAGGSHCLFNS
]		,		CCCCTSASLPARERHEWPGPAA
ł	}					AAAAAAAARRSQTQREGLFA
						GWGGGFAMSDDDSRASTSSSSS
	1					SSSNQQTEKETNTPKKKESKVS
						MSKNSKLL\SPRAKGIQKEL\AD
	Ì					ITLDPPPNCSAGPKGDNIYEWR
						STILGPPGSVYEGGVFFLDITFTP
						EYPFKPPKVTFRTRIYHCNINSQ
	1					\GVICLDILKDNWSPALTISKVL
						LSICSLLTDCNPADPLVGSIATQ
1						YMTNRAEHDRMARQWTKRYA
	1	1				Т

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23345	53713	Α	23479	372	1839	GEAGGTEAEPENGAGGPRVPG
		Ė				WWGWSSTSSPRGCWDSGRSAA
			:			AIMDFLWDKRTGLAARKMPHP
						RRYHSSERGSRGSYCEHYRS/R
						K/HKQRRSRSWSSSSDRTRRRR
İ						REDSYHVRRRCSRTFSRSSSQH
						SSRKAKSVEDDTEGHLIYHVGD
	1			İ		WLQERYEIVSTLGKGTFGRVV
	1					QCVDHRRRGARVALKIIKNVE
		l				KYKEAARLEIKVLEKINEKDPG
		l				KNLCVQMFDWFDYHGHMCISL
		1				ELLGLSTFDFLKDNNHLPYPIH
	•	1				QVHHMASQLCQAVKFLHDNK
		ŀ				LTHTDLKPENILFVNSDYELTY
		ļ				NLEKKRHERSVKSTAVRVGDF
						GSATFDHEHHSTIVSTRHYRAP
		ŀ				EVILELGWSQPCDVWSIGCIIFE
						YYVGFTLFQTHDNRQHLATME
						RILGPIPSRMIRKTRKQKYFYRG
						RLDWDENTSAGRYVRENCKPL
						RQYLTSEAEEDHQLFDLIESML
						EYEPAQRLTLGEALQHPFFSRL
		l				WAEPPNKLWDSSQDISP
23346	53714	Α	23480	28	941	QSARPSLKRARSQRGRPLPSRA
						LVFLFLFFFFFFNFLPIHTQPSAH
						KDMTTNAGPLHPYGPQHL\RLD
						NFVPN\DRPTWAYTGLGLFSVT
		1				GGLRSWTTWAVVQGRAAGCPI
						GDFGRATVPCCWVLQCVGFIH
						LVIEGLVPFSNYERPCFGDQSLL
						NLKLW\KEYGQGETARLHPGG
						DKFHSVALGTITSFA\WGEPFSL
					•	WVVIA\FLLQHPL\RFIL\QLVVS
						VG\QIYG\DVF\YFLKEHPRRDS
						KHGKPGPTPLLLPGFSFVFHEM
						PLWLVPAPGSPLCLDA\VKAPS
						THAPEHAWNAKGPQKPKSKKN

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide inscrtion)
				sequence		
23347	53715	A	23481	1	1386	MWRNLKLPRDLLNGFDQNADS
						DMDNEVQAEVVTDGDEELIGN
					·	WSKGHSCYILAKRLAAFFPCPK
						DLWKFELERDDLGYLAEEISSS
	•		ļ			EAFKSVSYEKAHAWNPLESTC
						RHASLALAVVLQRRDWENP\G
						VTQL\NRLAAHPPFASW\RNSEE
						ARTDRPSQQLRSLNGEWRLMR
						YFLLTHLCETLVKVKDAEDQL
						GARVCVLERQIFDFLGYQWAPI
						LANFVHIIIVILGLFGTIQYRPRY
						ITGILSFPKHQESLSLKFTCLTEA
						ERDTDALLHLENACRESTATQR
						IQGFFLKDKKRKKQQPQQQKQ
						QQWAKVQLGPWLQRVQAPSL
						GSFHGVLSLRVHRSKELRFRNC
						CLDFRGCMEMPGCPGRGVLQA
			.			QNPRGEPLLGQCRREMCGGSP
						YIVPTGALPSGAVRRRPPSSRPH
ļ						NGRPTDSLHCVPGKAADTQHQ
						PLKAARRAAVPCKATGSELPK
23348	53716	Α	23482	3	732	SPLTPTATSGNPSKASSPGGLGS
						LVSTASSLPAGEKNSPRTGPGP
						RRAG\AGIPNSGPDTVEGPYNL
			ļ			CGRQGRWGGLIISLASEMKLQT
						LQTSVTTAKATLLELFVPLGGL
						VASLASEVRLQIFPRGGHLQSA
						KCGLLLHFSLPTSRSGRRKANR
						PMAVTVLLVLIRDCADLLAVK
						AVIHCYQDQWPLRGSSRPANTP
			}			PPKASVSRRPPGHRGVRRGSPL
						LQVGARRVGAEAPALLELQSPS
23349	53717	A	23483	1	234	
23350	53718	Α	23484	1	256	MLNLQGKEWLELEEDAQKAYI
						MGLLDRLEVVSRERRLKVARA
						VLYLAQDGEKFCPKRDCTNNS/
						RY/NSI*MIQMMRFGTLELMRL
23351	53719	A	23485	1	3333	
23352	53720	A	23486	1	2661	
23353	53721	Α	23488	3	433	

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ł				sequence		
23354	53722	A	23489	3	990	EKMNTADQARVGPADDGPAPS
						GEEEGEGGGEAGGKEPAADAA
						PGPSAAFRLMVTRREPAVKLQ
						YAVSGLEPLAWSEDHRVSVST
						ARSIAVLELICDVHNPGQDLVIH
						RTSVPAPLNSCLLKVGSKTEVA
						ECKEKFAASKDPTVSQTFMLDR
						VFNPEGKALPPMRGFKYTSWSP
1						MGCDANGRCLLAALTMDNRL
1						TIQANLNRLQWVQLVDLTEIYG
						ERLYETSYRLSKNEAPEGNLGD
						FAEFQRRHSMQTPVRMEWSGI
					}	CTTQQVKHNNESRDVGSVLLA
					•	VLFENGNIAVWQFQLPFVGKES
		1		į		ISSCNTIESGITSPSVLFWWEYE
						HNNRKMSGLIVGSAFGPIKILPV
						1
						NLKAVKGYFTLRQPVILWKEM
						DQLPVHSIKCVPLYHPYQKCSC
		l				SLVVAARGSYVFWRLLLISKAG
						LNLHNSHVTGLHSLPIVSMTAD
						KQNGTVYTCSSDGKVRQVIPIF
						TDVALKFEHQLIKLSDVFGSVR
						THGIAVKPCGAYLAIITTEGMIN
	1	1				GLHPVNKNYQVQFVTLKTFEE
1						AAAQLLESSVQNLFKQVDLIDL
						VRWKILKDKHIPQFLQEALEKK
						IESSGVTYFWRFKLFLLRILYQS
						MQKTPSEALWKPTHEDSKILLV
						DSPG\MGNADDEQQEEGTSSKQ
						VVKQGLQERSKEGDVEEPTD\D
		l				SLPTTG\DAG\GREPMGRRNSW
		<u> </u>				GNSKGKIRSCWEMRLTREHMK
23355	53723	Α	23490	1218	1446	RCTDSQV\RSPTPLHLAAQACSL
						ETTVCLLCSKADYTLSEKRGW
						MPIHFAAFYDNVCIIIALCRKDP
						SLLEAEATAE
23356	53724	Α	23491	2	632	NMAKTYDYLFKLLLIGDSGVG
						KTCVL\FRFSE\DAFNSTFIFTIRI
		İ				DFKIRTIELDGKRIKLQIWDTAG
	1					QERFRTITTAYYRGAMGIMLVY
		1				DITNEKSFDNIRNWIRNTEEHAS
		1				ADVEKMILGNKCDVNDKRQVS
		1				KGS/GENKLALDYGIKFMETSA
						KANINVENAFFTLARDIKAKMD
		1				KKLDGNSP\QGSNQGVKITPDQ
1						QKRSSFFRCVLL
L		Ь	l		L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23357	53725	IA	23492	1	1018	MVFYHVGQAGLELLTSGPHCP
23337	33723		23472	1	1010	DSQAADKAESATLLSRARLGA
					<u> </u>	YCACASLERRWPHFPSGRECN
	į		1			MAKTYDYLFKLLL\IGDSGVGK
						TCVLFRFSEDAFNSTFISTIGIDF
						KIRTIELDGKRIK\LQIWDTAGQ
						ERFRTDHNRPTYRGAMGI\M\L
						VYDITNE\KSF\DNIRN\WI\RNIE
						EHASAD\VEK\MILGNKCDVND
						KRQVSKERGEKLALD\YGIKFM
						ETSAKANINVENAFFTLARDIK
						AKMDKKLGPARPHGTLLGKGE
						EFAGVAGASASGSASLGTLPKQ
						TRPCPVNPQASREHTAFSIPLKG
						QKDNMEGNSPQGSNQGVKITP
						DQQKRSSFFRCVLL
23358	53726	A	23493	1	130	
23359	53727	Α	23494	1	1392	
23360	53728	Α	23495	445	649	YPSLTNSKLLVHH\FMELC\WD
						KCVEKPQGIALDSRTENCLS\SC
						\VDRFIDT\TLAITSRFAQIVQKG
						GQ
23361	53729	Α	23496	210	397	VIYAPGKKAQRTQK/APKTKTK
						KPHLSPCPSP/WRSRSCQPMQQS
		ļ				WSGAQVHIPQPPHTTCITWA
23362	53730	Α	23497	55	286	QASSSPPISSVTCRKALPPGQTK
		l				KGLVASGDPEPVSMGTSSPTPS
		l		•		LAPEHSSRASLEPSSSA*PSLAT*
				:		SRAGGSQA
23363	53731	A	23498	780	1022	QAMHGRSPGVTVRASGLRNEG
						DGPLGSVPTQARATGAPGHSAT
						PSPAWRAW\SSSSKAS/WLPDA
		1				AGPPTPPWSGPSPPTSW
23364	53732	Ā	23499	1	782	
23365	53733	A	23500	275	688	
			1	<u>L </u>		

NO:	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23366 53734 A 23501 1166 2271	NO:			in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
23366 \$3734 A 23501 1166 2271 LRCILTLISRWCS*VSTVANLPP GISKWGANWQRCEMKE*CWSI AVTWCITCAQRSG/IGDSSPYCP ATSPNEYVKANLIWRGPVEQH PLVNYLDHERGTLSNFTARHYI PVLYVLRAILWARTPLPIPGGTI VSKVKMKKRILRAMRQKQQI PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAGFRMTQQDA PLVTAECITLESDWRSKEEVLK GMTDNLLLAGFRSTPHSK SEHHEQSTISVARLQAPVRWGD DEAQFIIMLTLNKHAAGDQHM RIFSTLARRIMHEEFRNALVA ASADAIASLLNMWTVKEHHE 23367 53735 A 23502 I 2604 23368 53736 A 23503 306 468 TRHCDRRQPRGHRRAGT/CNG I*GCCCVETKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AAQQGTMARSLLIPLQIILLLSL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHCGGVU VNERWLTAAHCKMNEYTVH LGSDTLZE/ADRKAQRIKPSKSF RHPGYFTOTHVNDLMLVKLNS QARLSSMVK 23370 53738 A 23505 I 1032 MMQATVGFEDGDAATSQVSPS VNKVHSSIYFLDPFEEYIVIAFK LRGPDISCSQIINGEDCSPHSQI WQAALVMEME/LFCSGVLVHP QWVLSAAPVSRKYNRPLLAND LMLKLDESVSESDTIRSISIASQ CPTAGNSCLVSGWGLLANDFT SRLIPPLECRNPVPQLLIPPEKI RTAPDTLGSSPYENSTGSSGS FLLPAPPSPFPRPHGTPPTSCG QDVDVPALAGGSLGSPAGEAL PRIAAPERLSRHIGKIPSPAPAI LTAILALETAGEEVPFPLTSCA WMSSSSPGLHEGLQCLTCKFI AVRWHPRLQEKRLQSVPSPDSF IPGARRPAAR	ł	sequence	ĺ	09/540,217	1 - 1	of peptide sequence	deletion, \=possible nucleotide insertion)
GISKWGANWQRCEMKE*CWSI AVTWCITCAQRSG/IGDSSPYEQH ATSFNEYVKANLTWRGPVEQH PLVNYLDHERGTLSNPTARHYI PVLYVLRALLWARTPLIPPEGTI VSKVKMKKRILRAMRQKQQQV PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAQFEMTQQDA PLVTAECITLESDWRSKEEVLK GMTDNLLLAGRCRYPRKLENQ LWAREAVFSTGLGFSFAIPHSK SEHIEQSTISVARLQAPVRWGD DEAQFIIMLTI.NKHAAGDQHM RIFSRLARRIMHEFERNALVNA ASADAIASLLNMNWTVKEHHE 23367 53735 A 23502 1 2604 23368 53736 A 23503 306 468 TRHCDRQPRRGHRRAGT/CNC 1*KGCCCVRTKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AAQQGTMARSILI.PLQILLI.SL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHGGGV VNERWVLTAAHCKMMEYTVH LGSDTL/E/ADRKAQRIKPSKSF RHPGYFTQTHVNDLMLVKLNS QARLSSMVK 1032 MMQATVGFEDGDAATSQVSPS VNKVHSSIYFLDFFEEVIVIALD QWVLSAAPVSRKYNRPLLAND LMLIKLDESVESDTIRSISIANS QCPTAGNSCLVSGWGLLANDFT SRUPPPLECRNPVPQLLLPERK RTAPDTLGSSPVPKSRTGSGSS FILPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLGSPAGES TLPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLGSPAGES TLPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLGSPAGES TLPAPPSFPRPHHGRIPSPAPAG LTAILALETAGEEVPFPLTSCA WMSSSSPPGLHEGLQCLTGKFI AVKWPRLQEKRLQSVPSPDSF IPGARRAAR					sequence		
GISKWGANWQRCEMKE*CWSI AVTWCITCAQRSGIGDSSPYEQH AISFNEYVARALIWARTPLIPEGTI TVSRVKMKKRILRAMRQKQQQV PLVVLRALIWARTPLIPEGTI VSRVKMKKRILRAMRQKQQQV PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAM CRTSLEVEHILLAQFRMTQQDA PLVTAECITLESDWRSKEEVLK GMTDNLLLAGRCRYPRKLENQ LWAREAVFSTGLGFSFAIPHSK SEHIEQSTISVARLQAPVRWGD DEAQFIIMLTINKHAAGDQHM RIFSRLARRIMHEFERNALVNA ASADAIASLLNMNWTVKEHHE 23367 53735 A 23502 1 2604 23368 53736 A 23503 306 468 TRHCDRQPRRGHRRAGTICNC 1*KGCCCVRTKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AAQGGTMARSLLPLQILLISL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHGGGV VNERWVLTAAHCKMEYTVH LGSDTL/E/ADRKAQRIKPSKSF RHPGYFTQTHVNDLMLVKLNS QARLSSMVK 1032 MMQATVGFEDGDAATSQVSPS VNKVHSSIYFLDFFEEVIVIALE QWVLSAAPVSRKYNRPLLAND LMLIKLDESVESDTIRSISLAND LMLIKLD	23366	53734	A	23501	1166	2271	LRCILTLISRWCS*VSTVANLPP
AVTWCITCAQRSG/IGDSSPYPC ATSFNEYVKANLTWRGPVEQD PLVNYLDHERGTLSNPTARHYI PVLYVLRAILWARTPLPIPGRTT VSKVKMKKRILRAMRQKQOYI PLLVGIGDEISMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHILLAQFRMTQQDA PLVTAECITLESDWRSKEEVLK GMTDNLLLAGRCRYPRKLEAD LWAREAVFSTGLGFSFAIPHISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTISVARLQAPVRISK SEHIEQSTRIVALAGEORIH RIFSRLARRIMHEFFRNALNNA ASADAIASLLMNNWTVKEHIE 23368 53736 A 23503 306 468 TRHCDRQPRRGHRRAGT/CNC PKCCCCVRTKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AQQGTMARSLLLPLQILLLSL ALETAGEAQGDKIDGAPCA GSHPWQVALLSGNOLHCGGVI VNERWVLTAAHCKMNEYTVH LGSDTLE/ADRKAQRIKPSKSF RHPGYPTOTHVNDLMLVKLNS QARLSSMVK QARLSSMVK LRRGPDISCSJINGEDCSPHSQI WQAALVMENE/LFCSGVLVHP QWVLSAAPVSRKYNRPLLAND LMILKLDESVSESDTIRSISIASQ CPTAGNSCLVSGWGLLANDPT SRLPPPLECRNPVPQLLLPPEK RTAPDTLGSSPVPKSRTGSSGSS FLLPAPPSPFRRHHGTPPTSCG QDVDVPALAGGSLGSPAGEAL PRLAAPERLSRHHGRIPSPAPAT LTAILALETAGEEVPFPLTSCA WMSSSSPPGLHEGLQGLTGKFF AVRWHPRLQEKRLQSVPSPDSR IPGARRAAR							GISKWGANWORCEMKE*CWSP
ATSFNEYVKANLTWRGPVEOH PLVNYLDHERGTLSNPTARHY PLLVYLRAILWARTPLPIPGRTI PVLVYLRAILWARTPLPIPGRTI VSKVKMKKRILRAMRQKQQVI PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAQFRMTQQDA PLVTABCITLESDWRSKEEVLK, GMTDNLLLAGRCRYPRKLEAD LWAREAVFSTGLGFSFAIPHSK SEHIEQSTISVARLQAPVRWGD DEAQFIIMLTINKHAAGDQHM RIFSRLARRIMHEEFRNALVNA ASADAIASLLNNWTVKEHHE 23367 53736 A 23502 1 2604 23368 53736 A 23503 306 468 TRHCDRRQPRRGHRRAGT/CNC I*KGCCCVRTKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AAQQGTMARSLLPLQILLLSL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHCGGVL VNERWLTAAHCKMMEYTVH LGSDTLÆ/ADRAKAQRIKPSKSF RHPGYFTQTHVNDLMLVKLNS QARLSSMVK MMQATVGFEDGDAATSQVSPS VNKVHSSIYFLDFEEEVIVIAFK LRRGPDISCSQIINGEDCSPHSQI WQAALVMENE/LFCSGVLVHP QWVLSAAPVSRKVNRPLLAND LMLIKLDESVSESDTIRSISIASQ CPTAGNSCLVSGWGLLANDPT SRLPPPLECRNPVPLLLIPEKS RTAPDTLGSSPVPKSRTGSSGSS FILIPAPPSPPRRHHGTPPTSCA WMSSSSPPGLHEGLQGLIGKFI AVRWHPRLQEKRLQSVPSPDSR IPGARRAAR							1
PLVNYLDHERGTLSNPTARHYI PVLYVLRAILWARTPLPIPGRTI VSKVKMKKRILRAMRQKQQV PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAGRCRYPRKLEAD LVAREAVFSTGLGFSFAIPHSK SEHHEQSTISVARLQAPVRWGD DEAQFIIMLTLNKHAAGDQHM RIFSRLARRIMHEEFRNALVNA ASADAIASLLNMNWTVKEHHE 23367 53735 A 23502 1 2604 23368 53736 A 23503 306 468 TRHCDRRQPRRGHRRAGYCNC PKGCCCVRTKQPLSSVLISPNF CPCSSFLKNS AQQGTMARSLLLPQILLLSL ALETAGEEAQQDKIDLGGGVL VNERWVLTAAHCKMNEYTVH LGSDTL/E/ADRKAQRIKPSKSF RHPGYFTOTHVNDLMLVKLNS QARLSSMVK 23370 53738 A 23505 1 1032 MMQATVGFEDGDAATSQVSPS VNKVMSSISYLLDFFEEVIVIAFK LRRGPDISCSQIINGEDCSPHSQI WQAALVMENELFCSGVUYHP QWVLSAAPVSRK YNRELLANDFT SRLPPPLECRNPVPQLLIPPEK RTAPDTLGSSPVFRSRTGSSGSS FLLPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLGSPAGEAL PRLAAPERLSRHHGKIPSPAPAL LTAILALETAGEEVPFPLTSCA WMSSSSPPGLHEGLQGLTGKFI AVRWHPRLQEEKRLQSVPSPDSI IPGARRPAAR				!			1
PVLYVLRAILWARTPLPIPGRTT VSKVKMKRILRAMRQKQQVI PLLVGLGLDELSMSAASPIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAQFRMTQQDA PLVTAECITLESDWRSKEVLK GMTDNLLLAGRCRYPRKLEAD LWAREAVFSTGLGFSFAIPHSK SEHIEQSTISVARLQAPVRWGD DEAQPIIMITLNHAAGDQHM RIFSRLARRIMHEEFRNALVNA ASADAIASLLNMNWTVKEHHE 23368 53736 A 23502 1 2604 23368 53736 A 23503 306 468 TRHCDRRQPRRGHRRAGTICNG I*KGCCCVRTKQPLSSVLISPNF CPCSSFLKNS 23369 53737 A 23504 102 509 AAQQGTMARSLLIPLQILLILSL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHCGGVL VNERWVLTAAHCKMNEYTVH LGSDTLÆJADRKAQRIKPSKSF RHPGYFTQTHVNDLMLVKLNS QARLSSMVK MMQATVGFEDGIDAATSQVSPS VNKVHSSIYFLDFFEEYIVIAFK LRQPDISCSQIINGEDCSPHSQI WQAALVMENE/LFCSGVLVHP QWVLSAAPVSRRYNRFLLAND LMLIKLDESVSESDTIRSISIASQ CPTAGNSCLVSGWGLLANDFT SRLYPPLECRNPYPQLLLPPEKS RTAPDTLGSSPVPKSRTGSSGSS FLLPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLSGAGEAL PRLAAPERLSRHHGKIPSPAPAC LTAILALETAGEEVPFPLTSCA WMSSSSPPGLHEGLQGLTGKFI AVRWHFRLQFEKRLQSVPSPDSI IPGARRPAAR							~
VSKVKMKKRILRAMRQKQQVI PLLVGLGLDELSMSAPSIPAAK ARMAQLDSRECRKLLNQAMA CRTSLEVEHLLAQFRMTQQDA PLVTAECITLESDWRSKEEVLK GMTDNLLLAGRCRYPRKLEAD LWAREAVFSTGLGFSFAIPISK SEHIEQSTISVARLQAPVRWGD DEAQFIIMLTUNKHAAGDQHM RIFSRLARRIMHEEFRNALVNA ASADAIASLLNMNWTVKEHHE 23367 53735 A 23502 I 2604 23368 53736 A 23503 306 468 TRHCDRQPRRGHRRAGTICNC CPCSSFLKNS 23369 53737 A 23504 102 509 AAQQGTMARSILLPLQILLLSL ALETAGEEAQGDKIIDGAPCAR GSHPWQVALLSGNQLHCGGVU VNERWVLTAAHCKMNEYTVH LGSDTLFE/ADRKAQRIKPSKSF RHPGYFTQTHVNDLMLVKLNS QARLSSMVK 23370 53738 A 23505 I 1032 MMQATVGFEDGDAATSQVSPS VNKVHSSIYFLDFFEEVIVIAFK LRGPDISCSQIINGEDCSPHSQI WQAALVMENEFLFCSGVLVHP QWVLSAAPVSRKYNRPLLAND LMLIKLDESVSESDTIRSISIASQ CPTAGNSCLVSGWGLLANDPT SRUPPPLECRNPVPQLLLPPEKF RTAPDTLGSSPVPKSRTGSSGSS FILPAPPSFPRPHHGTPPTSCG QDVDVPALAGGSLGSPAGEAL PRLAAPERLSRHHGKIPSPAPAE LTAILALETAGEEVPFPLTSCA WMSSSSPPGLHEGLQGLTGKFF AVRWHFRLQEKRLQSVPSPDSH IPGARRPAAR				,			1
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23371 53739 A 23506 303 393							IPGARRPAAR
	23371	53739	Α	23506	303	393	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23372	53740	A	23507	391	1170	GSFFFFFKYFAYSPMPKNSKVV
						KRELDDDVTESVKDLLSNEDA
						ADDAFKTSELIVDGQEEKDTDV
						EGK\SEVEDERPAWNSKLQYIL
						AQVGFSVGLGNVWRFPYLCQK
						NGGVVEPECEQSSATTYYWYR
						EALNISSSISESGGLNWKMTICL
						LAAWVMVCLAMIKGIQSSGKV
						NTYDRVTMSATVLALQEQWQT
						FSPIRSGNGHLTGSGAQRTPCRI
						RRGGSQLRVCDGGKQQWWTV
						SQSSARAITNTDQKSVQLQDLIE
23373	53741	Α	23508	45	546	EGGGSRSLWFRELIGQPVQSFIE
						PVAAGGAVALNVQ/RSCGVVS
						AAPACL*SRPHSWRWGGPACW
						RTPGG/LASRSSSSASILMSSSRA
						SFTRSLSLLSTHKDQGLVCSGSS
						GATGAGSC/LWPPTSHTVKLMF
						LYSTVSTLKPMVGMVVTISPSL
						SL*RIVVFPAASRPT/YEDAHFFF
						CQKGP*RGSWSSLQHPGPP
23374	53742	A	23509	1367	2142	VRRPQGCLGKRSLLSRVCALVP
						WGIWP/WSLAVGPQYSSLGSQP
						ILCASIPGLVPKQLRFCRNYVEI
						MPSVAEGIKIGIQECQHQFRGR
		l				RWNCTTVHDSLAIFGPVLDKG
						YFGYLETITVPSDSVVSTKAETP
						DKGVCGRELGFRGGWLPELQS
		1				LDEISMGWALCSRLQIKATRES
						AFVHAIASAGVAFAVTRSCAEG
						TAAICGCSSRHQGSPGKGWKW
						GGCSEDIEFGGMVSREFADARE
						NRPDARSAMNRHNNEAGRQV
			1			GSPPARVLGKKEPPQQGVCPGS
						LGHMARSLAVGPQYSSLGSQPI
						LCASIPGLVPKQLRFCRNYVEI
						MPSVAEGIKIGIQECQHQFRGR
						RWNCTTVHDSLAIFGPVLDKG
						YFGYLETITVPSDSVVSTKAETP
1						DKGVCGRELGFRGGWLPELQS
						LDEISMGWALCSRLQIKATRES
1						AFVHAIASAGVAFAVTRSCAEG
1						TAAICGCSSRHQGSPGKGWKW
						GGCSEDIEFGGMVSREFADARE
		[NRPDARSAMNRHNNEAGRQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23375	53743	Α	23510	2094	2619	ASPASPAPCSGIPRT*SPYPSPSR
			:			TQ\APRARGPAIHAHPGP/RGCT
						ALPRQGPPTPGSARAPPRIPAHR
			:			APTSPTLNFCLASSRQSMPPAPR
						PPPVPATNPVTPGCGRPPPTLGC
						RNPQTLRPPPGLPSPALLRRPLP
						PPARVGMPPHQPPLPTTPACWR
						PVDWARAEGAGRAVC
23376	53744	Α	23511	3050	3317	PARTSPSASGPP**DLISALWTPS
						QPRAPTRQSLNSLSSSAPSSSTIG
			ļ			PSPSPPAPGPAPPLPWLPPSPSSL
						PRLSQVSRRPHSHLHPQ
23377	53745	Α	23512	862	2954	AEHTLLVCFLNTSLTTTSTHVES
						FGGQLRGSQIGHPRRPNKMPYA
						WSIPQDPPLKPERDPGGPLGVS
						LSLPVNQYGESEEPMGDDDYD
						EYSKELNQYRRSKDSRGRGLSR
						GRGRGSRGRGKGMGRGRGRG
						GSRGGMNKGGMNDDEDFYDE
						DMGDGGGGSYRSRDHDKPHQ
						QSDKKGKVICKYFVEGRCTWL
						YHTTGNCINGDDCMFSHDPLTE
						ETRELLDKVEMTGSSVFDYIHP
						GDHSEVLEQLGLRTPTPGPPTPP
						SVSSSSSSSSSLADTPEIEASLTK
						VPPSSLVQERSFFVRMKSTLTK
]					RGLHVKASGYKVIHVTGRLRA
						HALGLVALGHTLPPAPLAELPL
						HGHMIVFRLSLGLTILACESRVS
						DHMDLGPSELVGRSCYQFVHG
						QDATRIRQSHVDCETHLHPPSL
						PTTPQTRASHSLVPGSPSSGLPT
			ļ			VPPAAPTAPGTPFSIGRAARRLS
						DASGLLDKGQVMTGYYRWLQ
						RAGGFVWLQSVATVAGSGKSP
						GEHHVLWVSHVLSQAEGGQTP
						LDAFQLPASVACEEASSPGPEPT
				•		EPEPPTEGKQAAPAENEAPQTQ
				:		GKRIKVEPGPRETKGSEDSGDE
				<u> </u>		DPSSHPATPRPEFTSVIRAGVLK
				}		QDPVRPWGLAPPGDPPPTLLHA
						GFLPPVVRGLCTPGTIRYGPAEL
						GLVYPHLQRLGPG\PALPEAFYP
						PLGLPYPGPAGTRLPRKGD
						GLVYPHLQRLGPG\PALPEAFY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			!	sequence		
23378	53746	A	23513	1319	2211	GSKVYGSLFHINHGNPFNMEVL
23378	33740	^	23313	1317	2211	VDSWPEYQMVIIRPQ/ETVRGE
	į					EKVQGNDLTFHHPTQEMTDDM
1						DSYTNVYRVFSKDPQKSQEVL
1						KNSEIINWKQKLQIQGFQESLG
İ						EGIRAAAFSNSVKVEHSRALLF
						VTEDILKLYATNKSKLGSWAET
						GHPDDELESETPNFKYAQLNVS
						YSGLVNDNWKLGMNKRSLRYI
						KRCLGALPAACMLGPEGVPVS
						WVTMDPSCEIGMGYSVEKYRR
						RGNGTRLIMRCMKYLCQKNIPF
						YGSVLEENQGVIRKTSALGFLE
22270	53747	ļ	22514	224	050	ASCQWHQWNCYPQNLVPL
23379	53747	Α	23514	234	858	SVPLQESPLPPPPTHQADC*SPC
						GAQGGWVTLAVASASS*AHTA
						SPLPRSPQAQGRAPSPAPPQIHL
						TTPSSFSSQGT/SHGGLGSTPGA
		1				DDPNTSSS\PPPGGSTPRWSAGD
						SAHSGGEVDGWS*SFP*SVPSY
1						GT*SPFCPWPSSGQRSFHGIASG
						WEPDGIPDGSSRWEPCGKPQES
						ETGVEGRQHFPKQLCLGSCRA
						VASPRVARA
23380	53748	A	23515	1	3582	DI DEGD A UDIVA CA DED DUIDIVI
23381	53749	Α	23516	628	4254	RLPESRAHDVAGADERPWPVH
		1				AGIFKSPAAGAAMSPAAAAAG
						AGERRRPIASVRDGRGRGCGGP
		1				AGAALLGLSLVGLLLYLVPAA
		1				AALAWLAVGTTAAWWGLSRE
		ļ				PRGSRPLSSFVQKARHRRTLFA
		1				SPPAKSTANGNLLEPRTLLEGP
		1		ł		DPAELLLMGSYLGKPGPPQPAP
				į.		APEGQDLRNRPGRRPPARPAPR
		1		i		STPPSPPTHRVHHFYPSLPTPLL
		}				RPSGRPSPRDRGTLPDRFVITPR
	ļ	<u> </u>				RRYPIHQTQYSCPGVL
23382	53750	Α	23517	1	392	KVHMNGLRYALYIMGPALTRG
						CHQVLIMLCCEVIERILT/VMM
						KEAYFGEVDIMDDLPT\GNISA
						NVTGRLYKCGVINHTVDLRRM
					1	KT/EKWQNNLLPSRQFGFIVLTT
		L				SAGIMDHEEARRKHTGGKILGF
23383	53751	Α	23518	227	352	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23384	53752	Α	23519	3	507	SFRAATMVRMNVLADALKSIN
						NAEKRGKRQVLIRPCSKVIVRF
						LTVMMKHGYIGEFEIIDDHRAG
						KIVVNLTGRLNKVGVQWRDLG
		ŀ				YCNLCLPGSSDSSASASLVAEIT
						\CGVISPRFDVQLKDLEKWQNN
	}					LLPSRQFGFIVLTTSAGIMDHEE
						ARRKHTGGKILGFFF
23385	53753	Α	23520	279	653	ARAQRFSSQVDVFYLIQSVIFCR
						LALAEFEIIDDPQSLGKLL*TLT
						GRLKQ/CLGVISPRF\DVQLKRP
ŀ						WKKWHEIIL/PSNPAKFGF\IVLT
ł						TSAGLHG/HHEGRQDRKHTGG
		1				KILGILLYRDVIHIFT
23386	53754	Α	23521	1	392	KVHMNGLRYALYIMGPALTRG
						CHQVLIMLCCEVIERILT/VMM
	1					KEAYFGEVDIMDDLPT\GNISA
						NVTGRLYKCGVINHTVDL/PPY
						ENGKWQNNLLPSRQFGFIVLTT
	}					SAGIMDHEEARRKHTGGKILGF
23387	53755	Α	23522	108	371	LSYIGKFEIIDDHRAGKIVVNLT
						GGLNKCGVISLRFDVQLKDLEK
		1				WQNNLLPSCQFGFIALTTSAGI
						MDHKEAR*KHTAGKILEFFF
23388	53756	Α	23523	252	378	
23389	53757	Α	23524	3	518	HASAFRAATMVRMNVLADAL
						KSINNAEKRGKRQVLIRPCSKVI
i						VRFLTVMMKHGYIGEFEIIDDH
				1		RAGKIVVNLTGRLNKVGVQWR
		1				DLGYCNLCLPGSSDSSASASLV
1		1				AEIT\CGVISPRFDVQLKDLEK\
		l				WQNNLLPSRQFGFIVLTTSAGI
						MDHEEARRKHTGGKILGFFF
23390	53758	Α	23525	279	709	ARAQRFSSQVDVFYLIQSVIFCR
1						LALAEF*NPLMTHQSLGKFVVN
		1				PHRAGLNKCG\VISPQILTVQLQ
		ł				RPWKKWAGLILLP\SRQVLGF\I
						VLTTSAGI\MDHEEARPKTPRR
						GKSWGFFFLGDVNYILFYKLKC
l		<u>L</u> _				LMDSGASTWSF
23391	53759	Α	23526	207	564	SMAKEAISINRSLWAGPAKISSP
		1	1			AETE\IESLISLFQKYVGKGGYN
						CTLSKTEFLSFMNAELAAFTKN
		1				QKDPGVLHRMMKKLGTNNDG
						QLDFSEFLNLIGGLAMACHDSF
						LKAVPSQKRT
						-

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23392	53760	A	23527	1	407	FREKQSQELSQYGAMGAGQSF
				1		NSQFLQHGGPRGPSVPAGMNP
			1			TGIGGVMGPSGLSPLAMNPTRA
						AGMTPLYAGQRLPQHGYPGPP
						QAQP/PAPTGGQENLL*GVSRA
		ŀ				AVSARSLPGPFRAIPVSHLAGNP
						ТРРМПРН
23393	53761	Α	23528	3	315	
23394	53762	Α	23529	1	763	DYEEITIDPTCSWKPVPVKPDM
						HIKEEPDGPALKRCRTVSPAHV
	}	1				LMPSVMEMIAALGPGAAPFAPL
			İ			QPPSVPAPSDYPGQGSSFLGPGT
ŀ				İ		FPESFPPTTPSTPTLAEFTPGPPPI
ĺ		ł				CYQSDIPSSLLTSEKST/SLPLQA
						RR/PPAGHLDPTHNPGTPGLHTS
						NLGAPPGPQLHHSNPPPASRQS
						LGQASLGPTGELAFSPATGVMG
			ļ			PPSMSGAGEAPEPALDLLPELT
						NPDELLSYLGPPDLPTNNNDDL
						LSLFENN
23395	53763	Α	23530	1	2901	MERRGPGAATARGRARPGGGP
						SVGLLATGSSLNPSFHGVARIVP
						GFIRIARPRDGSFAYESVPWQQS
						ATQPAGSLSVVTTVWGVGNAT
						QSQVLGNPMGPAGSPSGSSMM
	1					PGVAGGSSALTSPQCLGQQAFA
						EGGANKGYVQQGVYSRGGYP
						GAPGFTTGYAGGPGGLGLPSH
:						AARPSTDFTQAAAAAAVAAAA
						ATATATATVAALQEKQSQE
						LSQYGAMGAGQSFNSQFLQHG
						GPRGPSVPAGMNPTGIGGVM
23396	53764	Α	23531	80	412	KPGNGACAGREWCDGGGAAW
						NWRDPGLPVGDSGVWDRVLLE
						LLGPRSPRPLDVGGPAAGTPGV
				1		LSR/HLPVDCGLGPEALLCSSAP
						AARRPSLCGGHGRVMRGFRLG
						CGFRP
23397	53765	Α	23532	1	192	RPLDVGGPAAGTPGVLSR/HLP
						VDCGLGPEALLCSSAPAARRPS
1						LCGGHGRVMRGFRLGCGFRP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23398	53766	A	23533		2094	RPLDVGGPAAGTPGVLSRPCPS TAALAPKPFCAAPRPQPDAPAC AGATGGSCADFDSGVDFVRRR SSGLWGPQPPLFPVKNYTEMFQ DPVAFKDVAVNFTQEEWALLD ISQKNLYREVMLDTFWNLTSIG KKWKDQNIEYEYQNPRRNFRS VTEEKVNEIKEDSHCGETFTPV PDDRLNFQKKKASPEVKSCDSF VCGEVGLGNSSFNMNIRGDIGH KAYEYQEYGPKPCKCQQPKKA FRYRPSFRTQERDHTGEKPYAL LKNDGKKPLFYHSSIQRHMVV HSGDGPYKCKFCGKAFHWLSL YLIHERTHTGEKPYECKQCGKS
						FSYSATHRIHERTHIGEKPYECQ ECGKAFHSPRSCHRHERSHMGE KAYQCKECGKAFMCPRYVRRH ERTHSRKKLYECKQCGKALSSL TSFQTHIR\MHSGERPYECKTCG KGFYSAKSFQRHEKTHSGEKPY KCKQCGKAFTRSGSFRYHERTH TGEKPYECKLYGKALSRLISFRRH MRMHTGERPHKCKICGKAF\SS PSSFQRHERSHTGEKPYKCKQC GKAFTCFHF\FQYHERTHTGEK PDGCKQCGKAFRSAKYIRIHGR THTGEKPYECKQCGKAFHWVS SFHRHERTHAGEKPYECKHCG KAFTCSIYIRIHERIHTGEKPYQ CKECGKAFIRSSYCRKHERTHTI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23399	53767	A	23534	48	1317	CGLMDELVHDLTSALEQTSEQ
						NKLGELWEEMALSPROORROL
ļ						RKRRGRKRRSDFTHLAEHTCC
						YSEASESSLDEATKDCREVAPV
						TNFSDSDDTMVAKRHPALNAI
		1				VKSKQHSWHESDSFTENAPCRP
						`
						LRRRR\KVKRVTSEVAASLQQK
						LKVSDWSYERGCRFKSAKKQR
		ļ				LSRWKENTPWTSSGHGLCESA
						ENRTFLSKTGRKERMECETDEQ
		ŀ				KQGSDENMSECETSSVCSSSDT
						GLFTNDEGRQGDDEQSDWFYE
		İ	İ			GECVPGFTVPNLLPKWAPDHCS
						EVERMDSGLDKFSDSTFLLPSR
		ł				PAQRGYHTRLNRLPGAAARCL
		l				RKGRRRLVGKETSINTLGTERIS
]						HIISDPRQKDFWLPSAGKRERN
İ		1				QFNPLSPLYSLDVLADASHRRC
	İ					SPAHCSARQANVHWGPPCSRDI
						KQEAETSGHSIFV
23400	53768	Α	23535	225	403	
23401	53769	Α	23536	1	403	REQEFLHMAPGEVTITVR\LIRS
						LEHR\NFKPVVYHGVNLDQTV
		l				KEFIVFLKQDIPLRTNLPPPFRN
		1				YKYDALKIIHQAHKSKTNELVL
		1				SLEDDERLLLKEDSTLKAAGIA
						SETEIGIF\REEDYRNYKANPISS
23402	53770	A	23537	2	239	
23403	53771	A	23538	2	214	
23404	53772	Α	23539	1	1341	
23405	53773	A	23540	1010	1677	
23406	53774 53775	A A	23541	1010	1509	
23407				223	6355	VIANDICENED DEFERIENTS CNIC
23408	53776	Α	23543	223	10504	VKMPIGSKERPTFFEIFKTRCNK
	1					ADLGPISLNWFEELSSEAPPYNS
						EPAEESEHKNNNYEPNLFKTPQ
						RKPSYNQLASTPIIFKEQGLTLP
		ŀ				LYQSPVKELDKFKLDLGRNVP
		İ		ļ	ĺ	NSRHKSLRTVKTKMDQADDVS
						CPLLNSCLS\ESPVVLQ\CTHVTP
						QRDKSVVCGSLFHTPKFVKGR
						QTPKHISESLGAEVDPDMSWSS
						SLATPPTLSSTVLIVRNEEASET
						VFPHDTTANVKSYFSNHDESLK
L	<u></u>					KNDRFIASVT
23409	53777	В	23544	137	709	
23410	53778	Α	23545	1	1974	

SEQ ID NO:	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NU:	sequence	noa	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23411	53779	Α	23546	207	1622	MACCL\SDEVKESKRINAEIEKA
						AAAETRRDARREL/RRLLLLGT
						GESGKSTFIK\QMRIIHGAGYSE
						EDKRGFTKLVYQNIFTAMQAM
						IRAM\ETL\KILYKVRSRTRANA
		ŀ		-		L\LIRGGGTLEKVTTFGAFSYVQ
						LPLKDLFGRDPGIQE/CAYERQP
						RSYQLSDSAKYYLTDVDRIATL
						GYLPTQQ\DVPAGSRVPTT\GIIE
						YPFDLENIIF\RMVDVGG\QRS\E
						RRKWIHCFENVTSIMFLVALSE
'						YDQVLVESDNENRMEESKALF
						RTIITYPWF\QNSSVILFLNKKDL
						LEDKI\LYSHLVDYFPRSSMGPQ
		1				R\DAKAAREFILKMFVDLNPNS
						DKII/YTSHFTCPHRHRRTSRFVF
						PAVKDTILQLNLEEYKPGLSAP
						GPGRRDGDTGQDLPSTEPAAA
		1				RAGGAAESGPGPLPAGGDFFFF
						IFLTNGFYFTVIRGCTSLPPYTSR
						TFSPFVNGKGSLFLALTYGSLFS
						KKKKKKKE
23412	53780	Α	23547	1	401	KKVPGRLSEMSFSLNFTLPANT
						TSSPVT\DCGPSLGLAAGIPLLV
						ATALLVALLFTLIHRRRSSIEAM
						EESDRPCEISEIDDNPKISENPRR
						SPTHEKNTMGAQEAHIYVKTV
	ļ	4.		1.0	1051	AGSEEPVHDRYRPTIEMERRR
23413	53781	Α	23548	584	1071	PCRLPLHLILVEDLSFSLNFTLP
						ANTVSTAAPIHQAT/PEKTSAEF
						TDCGPSLGLAAGIPLLVATALL
			1			VALLFTLIHRRRSSIEAMEESDR
						PCEISEIDDNPKISENPRRSPTHE
						KNTMGAQEAHIYVKTVAGSEE
						PVHDRYRPTIEMERRRGLWWL VPRLSLE
23414	53782	A	23549	25	654	LFLQMKDTSPIIFVRKELTIRQC
23414	33/62	A	23349	23	034	TKKVPGRLSEMSFS\LNFTLPAN
						TTSSPVTGGKETDCGPSLGLAA
						GIPLLVATALLVALLFTLIHRRR
						SSIEAMEESDRPCEISEIDDNPKI
						SENPRRSPTHEKNTMGAQEAHI
						Y\VKTVT\KSEEPVHDRYRPTIE
			1	1		MERRRGLWWLVPRLSLGMMP
						TSNQGSHQTWHLGTGLKTQGF
	}					VLGEERCQAAS
23415	53783	A	23550	108	413	RRSGSMALWRAYQRALAAHP
	33703	' '		1.00		WKVQVLTA/GLLGRKAALLAP
		1				CQ*LYLPTQASWTEGHASCSSL
						QEATSCPSAFCA*NSKATKQLV
						ERRGLQEHQRGRTLTMVS
L	l		L	<u></u>	<u>. </u>	S. C. C. D. C. C. C. C. C. C. C. C. C. C. C. C. C.

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23416	53784	Α	23551	Į.	146	
23417	53785	Α	23552	2	442	GREARRSRSMALWRAYQRALA
	}					AHPWKVQVLTA\ASGEEGSTA
						GPMPVAIPADPGLLD*RTRFLLL
						TPRSHFLPLCLLRMKLKSHKRS
						LMGLGDIISQQLVERRGLQEHQ
	}]			RGRTLTMVSLGCGFVGPVVGG
						WYKVLDRFIPGTTKVDA
23418	53786	Α	23553	3	675	RHSRFGAQGSMALWRAYQ\RT
						LF\ALPWK\VQVLTAG\$LMGLG
						DIISQQLVERRGLQEHQRGRTL
						E/TMVSLGCGFVGPVVGGWYK
			1			VLDRFIPGTTKVDALKKMLLD
						QGGFAPSFLGCFLPLVGALNGL
						SAQDNWGQTTAGELG/RDYPD
						ALITNYYLWPAVQ\LANFYLVP
			ŀ			LHYRV\AVVQCVAVIWNSYLS
			1			WKAHRALSLPHSIVSTLQVMQ
						LDPWNGSDNPPQKWA
23419	53787	Α	23554	512	1227	LQGHLMGIMAASRPLSRFWEW
						GKNIVCVGRNYADHVREMRSA
						/VWLSEPVLFLKPSTGVRAQRA
						RPILMPAYTRNLHHELEL\ALVI
						GQ\RCRAVPE\AAAMDYVGG\Y
			•			ALCL\DMTAWDVQ\ASARKKG
						LAWDSGE/DAFTASCPVQRVSL
			<u> </u>			PKEKIPDPSQ\LKALGFKVNGEL
					İ	RQEGETSSMIF\SIPYIISYVSKIIT
						LEEGDIILTGTPKGVGPVKEND
		L				EIEAGHTRAGQYDHLKVEKAQ
23420	53788	Α	23555	192	436	RTSDHECFREGQICSVCLRERL
						NLNPEEACLLSQAPCMLFVLQY
						LNTVIPYEKKASPPSV\EDLQML
L						\TNILFAMKEDNEKVP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23421	53789	Α	23556	94	1252	ARMEAPLVSLDEEFEDLRPSCS
						EDPEEKPQCFYGSSPHHLEDPSL
	,					SELENFSSEIISFKSMEDLVNEF
						DEKLNVCFRNYNAKTENLAPV
						KNQLQIQEEEETLQDEEVWDA
:						LTDNYIPSLSEDWRDPNIEALN
						GNCSDTEIHEKEEEEFNEKSEN
						DSGINEEPLLTADQVIEEIEEMM
						QNSPDPEEEEEVLEEEDGGETSS
		ĺ				QADSVLLQEMQALTQTFNNNW
						SYEGLRHMSGSELTELLDQVEG
						AIRDFWE\ELVQQLARRDELEF
,						EKEVKNSFF\TVL\IEVQNKQK\E
						QRELMKKRRKEKGLSLQSSRIE
						KGNQMP\LKRFSMEGISNILQSG
						IRQTFGSSGTDKQYLNTVIPYEK
						K\AYPPSVE\DLQMLTNILFA\M
						KEDNEKVPYFA
23422	53790	Α	23557	391	1169	GSFFFFFKYFAYSPMPKNSKVV
		1				KRELDDDVTESVKDLLSNEDA
		[ADDAFKTSELIVDGQEEKDTDV
		l				EGK\SEVEDERPAWNSKLQYIL
						AQVGFSVGLGNVWRFPYLCQK
						NGGVVEPECEQSSATTYYWYR
						EALNISSSISESGGLNWKMTICL
						LAAWVMVCLAMIKGIQSSGKV
						NTYDRVTMSATVLALQEQWQT
						FSPIRSGNGHLTGSGAQRTPCRI
						RRGGSQLRVCDGGKQQWWTV
		<u> </u>			7.0	SQSSARAITNTDQKSVQLQDLIE
23423	53791	A	23558	321	568	CLWCGASWPRRHSSPGRPAGD
		1				HRGDRP/EGPDTQGPGRPEPSTE
						PAFNPQSSLDQGGGPAAPAVD
22.42.4	62,700	_	22550	7.0	(00	DDTRRGHGGVCGDGGAGEA
23424	53792	Α	23559	510	690	LRSPQ\RGVAAPPFLGSPVPFGG
						FRVQGGMAGQPSAPPALQTDD
22425	52702	_	22560	25	490	PYSKAHSVTPKSTPRVT
23425	53793	Α	23560	25	480	VD A VOCI HEHEDWID //CORSE
23426	53794	Α	23561	84	292	VRAVQSLHFHFPWRR/ISCPSES TRSIHVDRLPRRNSASLSLSAFP
						VSAGALINGEAWVRMGQVPIP
	<u> </u>		<u> </u>		L	KAP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		İ
23427	53795	A	23562	1138	1924	CRSSATWTPWTTTGCRGDLTK
23 127	33775	-	20002	50		TYSLEAYDNWFNCLSMLVATE
		1				VCRLLLSPGSAMLRSPATFQVV
		ŀ				KKKHRTRMLEFFIDVARECFNI
		1				GNFNSMMAIISGMNLSPVARLK
		1				KTWSKVKTAKFDVLEHHMDPS
		ł				SNFCNYRTALQGATQRSQMAN
		İ				SSREKIVIPVFNLFVKDIYFLPQ
						NP\SNHLPNGHINFKKFWEISRQ
						IHEFMTWTQVECPFEKDKKIP\S
		Ì				YLLTAPHPTARKLSSSPSFESEG
		1				PENHMEKDSWKTLRTTLLNRA
23428	53796	A	23563	3	260	
23429	53797	Α	23564	1	1386	
23430	53798	Α	23565	137	611	MKVNLGDSQNLACVFCRKHD
		ŀ				DCPNKYGEKKTKEKWNLTVH
		1				YYCLLMSSGIWQRGKEEEGVY
		1				GFLIEDIRKEVNRASKLKCCVC
			İ			KKNGAFIGCVAPRCKRSY\HFP
						CGLQRECIFQFTGNFASFCWDH
						RPVQIITSNNYRESLPWHHLLG
						NLFEPIPSY
23431	53799	A	23566	181	486	DPQGEAKGAAPAEGGGGPV*W
		İ				N\GLQGPAGVPGRDGSPGANGI
				-		PGTPGIPGRDGFKGEKGECLRE
]	1				SFEESWTPNYKQCSWSSLNYGI
	<u> </u>		L			DLGKIAECTFTKMRS
23432	53800	Α	23567	600	888	QYNGMCSPGPAGVPGR\EGSPG
				ŀ		ANGIPGTPGIPGRDGFKGEKGE
		İ				CLRESFEESWTPNYKQCSWSSL
						NYGIDLGKIAGRNCWHILAPVP
			ļ			NIVSLYFGS
23433	53801	Α	23568	91	1170	ISLHRRPGPTCSMESPASSQPAS
		Ì				MPQSKGKSKRKKDLRISCMSKP
		1				PAPNPTPPRNLDSRTFITIGDRN
						FEVEADDLVTISELGRGAYGVV
						EKVRHAQSGTIMAVKRIRATV
						NSQEQKRLLMDLDINMRTVDC
						FYTVTFYGALFREGDVWICME
						LMDTSLDKFYRKVLDKNMTIP
						EDILGEIAVSIVRALEHLHSKLS
				-		VIHRDVKPSNVLINKEGHVKM
						CDFGISGYLVDSVAKTIN\AGCK
						VKHGHESLN\PDLNRKGCDVKS
						HVWSLGITMIEMAFLRFPYESW
						GTPFQQLKQVVEHPSPQLPAGR
						FSPEFVDFTAQCLRKNPAERMS
						YLELMEHPFFTLHKTKETDIA\A
						FVKEILGEDS
23434	53802	В	23569	111	193	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23435	53803	A	23570	312	696	RP*SAHPEPGDCPSPPHPPRCGG VAQTAQVPDAPAPV*AG\QHSR SYMEDHLKNKNRLEKEWEALC AYQAEPNSSFVAQREENVPKN RSLAVLTSQPTRNQCLPFLQHG PGDICTPVSISSIVLCHRVT
23436	53804	A	23571		2974	QAAGMGPPLPLLLLLLLLPPR VLPAAPSSVPRGRQLPGRLGCL LEEGLCGASEACVNDGVFGRC QKVPAMDFYRYEVSPVALQRL RVALQKLSGTGFTWQDDYTQY VMDQELADLPKTYLRRPEASSP ARPSKHSVGSERRYSREGGAAL ANALRRHLPFLEALSQAPASDV LARTHTAQDRPPAEGDDRFSES ILTYVAHTSALTYPPGPRTQLH EDLLPRTLGQLQPDELSPKVDS GVDRHHLMAALSAYAA
23437	53805	A	23572		3202	ASRRLPRGHGLSAAGQAAGMG PPLPLLLLLLLLPPRVLPAAPSS VPRGRQLPGRLGCLLEEGLCGA SEACVNDGVFGRCQKVPAMDF YRYEVSPVALQRLRVALQKLS GTGFTWQDDYTQYVMDQELA DLPKTYLRRPEASSPARPSKHS VGSERRYSREGGAALANALRR HLPFLEALSQAPASDVLARTHT AQDRPPAEGDDRFSESILTYVA HTSALTYPPGSRTQLREDLLPR TLGQLQPDELSPKVDSG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23438	53806	Α	23573	1	1459	MPLTSQDVDQKTAAWVPTNRD
						STVPSDPWDCKRRPWPLLVLRP
1					;	LAPGTKAPPGGQNGRGRQKGS
					}	PCSRRLLLEPAGPLGPASVRCIR
						TTTPQLLIITTSLRFPSLEAAVGY
						GSNTCIFANACQMVFIMITWPY
						LQCIPYSVLLKDLEMRNLRELE
						DLIIEAVYTDIIQGKLDQRNQLL
		١.	}			EVDFCIGRDIRKKDINNIVKTLH
			1			EWCDGCEA\ALLGIEQQVLRAN
						QYKENHNR/TLQQQVEAEDVYI
						LPQSGNDCRGLRFVSSLIGGVR
ļ						CVHLKHKGDAVVWEAIEEGEA
						LPGKYRLKGGRSVWAAAVRA
						VCMDQVTNIKKTLKATASSSA
						QEMEQQLAERECPPHAEQRQP
						TKKMSKVKGLVSSSPLGPAGA
						AGTHQAWVRWGGDTKGPFPPL
						STCSEFOTCPSPHQRLPTLLVLF
						QKNCYSPSPTPSFPSCSLQTQGL
						1 1
						HQCHPKTGSDTAQLPSRRFLSL
23439	53807	A	23574	2	396	CKGLVSLPVFLLLPRHFVRLVIK
23440	53808	A	23575	1	1014	
23441	53809	A	23576	110	212	
23442	53810	A	23577	1	747	
23443	53811	A	23578	323	495	GRAGKWHTQPE*SLRKAM*MF
						QILMMKMIVALARATSGTEAV
	,					AGRSWASGLRRSGRV
23444	53812	A	23579	26	229	
23445	53813	Α	23580	1	574	
23446	53814	A	23581	889	1189	
23447	53815	Α	23582	170	625	GAAVALVRPNGGTAASHVVCR
						EMRPLRPLQSRRCSRRRTGISN
						MRALENDFFNSPPRKTVRFGGT
						VTEVLLKYKKGETNDFELLKN
						QLLDPDIKDD\QIINW\LLEFRSS
						VMYLTK\DFE\QLISIILRLP\WL
						NRSQ\TVVEEYLAFLGNLVSA
23448	53816	Α	23583	166	455	RWKPSGTWPRNPMERPGH*ES
Ì						VPKPGD*AKKSGGET\RGRKEG
					}	NEHRETFSQIPDCHLNKKSQTG
E 			1			VKPCKCSVCGKVFLRHSFLDRH
İ			1			MRAHAGHKRS
					i	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23450	53818	A	23585	1	1970	MVTDRKAAAGTHGEARPCWG
25450	35010	``	25505	1		SGSRVHTYRGAGERAMRVHRP
						MPREDSERVREADGGKHEQRS
			i			FGDLHPTAAPAPAASGCGTQPG
						RVKQPGEWQHREAPQQRQPPV
						RPHGSHQLVTVRYERRPGGKE
						1
						RARLRPGLGCPSGRTPQGRAPQ
	1					GTALPTAAAAAIFPSAALTSGQ
						RNASLPKFYFRGGTWGTSQSEA
						LCGKLSNQFLGEPLVPRIRLILS
						CGTCSGCSSPRKDPGHPEAGNG
						PSRDFFQLPSEERDSVAFEDVA
						VNFTQEEWALLSPSQKNLYRD
						VTLETFRNLASVGIQWKDQDIE
		1				NLYQNLGIKLRSLVERLCGRKE
	1					GNEHRETFSQIPDCHLNKKSQT
						GVKPCKCSVCGKVFLRHSFLDR
		1				HMRAHAGHKRSECGGEWRETP
		1				RKQKQHGKASISPSSGARRTVT
						PTSMRPYE\TKVCGKA\FIS\PIYF
İ						QIHQRTPTEKRSYKCREIVRAFT
		1		1		VSSFFRKHGKMHTGEKRYECK
						YCGKPIDYPSLFQIHVRTHTGE
						KPYKCKQFGKAFISAGYLRTHE
						IRSHALEKSHQCQECGKKLS\CS
						SSLHRHERTHSGGKLYECQKC
						AKVFRCPTSLQAHERAHTGERP
		1				YECNKCGKTFNYPSCFRRHKKT
		1				HSGEKPYECTRCGKAFGWCSSL
		1				RRHEMTHTGEKPFDCKQWVKS
23451	53819	Α	23586	86	390	LGSGDLPWEINPLSSCSLLREKD
		1				PPTTSGPQT\TSPRNISPISNPELA
		İ				THAGNLATGPRNARSPGFLLSR
		ŀ				VPSVWDPTENRTVQLTWQPLP
						ELLELWPKVL
23452	53820	Α	23587	1	502	MLLTQSLFGGLFTRTHMKFGA
				•		VTQIRGPPLGDKSPVLLLFALER
						QRRHVLSMDPKLRCWSRTGKA
						AFPWCLIIAEMPDYSPTFQRCQ
		1				TTQGRLPWSFTLSSKSRFSGEG
						ARACYKCQKSDHQAKECPQPG
		į				IPPKPCPICAGP\TGNRTVQLTW
İ						QPLPEPLELWPKAL
23453	53821	В	23588	1	663	
23454	53822	Α	23589	233	496	
23455	53823	Α	23590	2	378	ERQRRHVLSMDPKLRCWSRTG
						KAAFPWCLIIAEMPDYSPTFQR
						CQTTQGRLPWSFTLSSKSRFSG
						EGARACYKCQKSDHQAKECPQ
						PGIPPKPCPICAGP\TGNRTVQLT
1		1				WQPLPEPLELWPKAL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23456	53824	A	23591	3	452	VRPRRDACLGPSPLAASPAFLG KGQVPQPLISLCPDPLFPHPNLIS LRPNPLCPHPDLVSLCPDPFPAF LEAHKNFQTPEPQQPGIPPEPPP PGACYKC/HEI*PPGQGMPAAQ DSS*AMS/LSVRDPTGNQTVQL TWQPLPEPLELWPKAL
23457	53825	Α	23592	1352	1540	
23458	53826	Α	23593	177	383	EAMEPWKGVPLGRVNTEQIAP FSEAEGPREIH*GSRKVRKAAQ GLNEGPRELFWQPLPEPLELWP KAL
23459	53827	Α	23594	1478	1744	
23460	53828	A	23595	23	178	SCLEVCDEQGPEK\TRQRALRG VSSVTEDTLNICRLCWQPLPEPL ELWPKAL
23461	53829	Α	23596	1	282	
23462	53830	A	23597	34	241	SGDLPWEINPPSSCSLLREKDPP MTSGPQT\TSPRNISPISNPDPTG NRTVQLTWQPLPEPLELLPKAL
23463	53831	Α	23598	121	286	AASFYSSPTSLTIPQPLSPFNLGV TLQSLPSLNFNSF\LFWQPLPEPL ELWPKAL
23464	53832	A	23599	144	488	GYEIWCRDSDQGTFPWEINPPS SCSLLREKDPPTTSGPQT\TRPR NISPISNLVSGLFLLSSPPSLTIPQ PLSPFNLGATLQSLPSLNFNSFH FLVETKETRFIRGAKIPAPVTD
23465	53833	Α	23600	471	677	
23466	53834	В	23601	67	555	
23467	53835	Α	23602	1	1440	
23468	53836	A	23603	860	1209	IPGNCRDSSGVGIKERETNAGS QHMH/IKNSSVHGYPRLCVEDE DAYKKQFSQYIKNSVTPDMME EMYKKAHAAVPENPVYEKKPK KEVKKKRWNRSKMSLAQKKD WVAQKKASFLRA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23469	53837	A	23604	1	966	MGFVKVVKNKAYFKRYQVKF
<u> </u>						RRRREGKTDYYARKRLVIQDK
						NKYNTPKYRMIVRVTNRD\IICQ
						IANA\RIEG\DMIVCAP\YAHELP
						KIWV*RVGLTNYAAA\YCTGLL
						AGPAGFFHRFGMDQIYE\GQV\E
						LTGDEYNVESIDGQPGCLHPAY
						WDARPLPRTYPLAIKVFGCP*R
						GA\VDGGLVLFP\HSTKTIPPGY
						DS\ESK\EFNAES/IHRKPHPWAQ
						NVARLHAPTLMEEDEDA/YTK
					:	KQFSQYIKNSVTPDMMEEMV/Y
						RKAHAAIRENP\VYEKKPKKEV\
						KRKRWNRPKMSLAQKKDRVA
						PKKASFLQSSRSGLLESLTPAIFP
23470	53838	A	23605	1	644	ANSSPVNPVVFFDVSIGGQEVG
						RMKIELFADVVPKTAENFRQFC
						TGEF/RVSFRCV*LLCLLQRGVL
						GLQ*GYR*TTCWP*MIAGDSDR
						RKDGVPIGYKG\STFH\RVIKDF
						MIQGGDFVNGDGTGVASIYRG
				:		PFADENFKLRHSAPGLLSMANS
						GPSTNGCQFFITCSKCDWLDGK
						HVVFGKIIDGLLVMRKIENVPT
						GPNNKPKLPVVISQCGEM
23471	53839	Α	23606	638	1391	CPAYKGAHGIQCVYCTLPPPLL
						QTTRRVRRNPTLPTSASGSEPW
						RLANSSPVNPVVFFDVSIGGQE
						VGRMKIELFADVVPNTAENFR
						QFCTGEFRKDGVPIGYKGSTFH
						RVIKDFMIQGGDFVNGDGTGV
						ASIYRGPFADENFKLRHSAPGL
						LSMLTWPGKLTAGIFGFHQANS
						GPSTNGCQFFITCSKCDWLDGK
						HVVFGKIIDGLLVMRKIE\GMPC
						IFQPTWTHFLPVITTACSALGSM
		_				LAAERGTKN
23472	53840	Α	23607	2	591	ARAEFGTSRVGAMAVANSSPV
		1				NP\VVFFDVSIGGQEVGRMKIEL
						FADVVPKTAENFRQFCTGEFRK
						DGVPIGYKGSTF\HRVIKGF\MI\
		ĺ		1		QGGDFV\NGDGTGSRPVFTRGP
						ILQMKIFKL\RHSAP\GLLSMAN
						SGP\STNGC\QFFITC\SKCDW/V
				1		WDGKHVVFGK\IIDGLLVMRKI
				<u> </u>		E\NVPTGPNNKP\KLPVVISQCG
23473	53841	Α	23608	6	424	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23474	53842	Α	23609	3	470	DAWVSGRLTELPSSCFFPPIFRQ
[1		LLPVMPEPSKSAPAPKKGSKKA
		ŀ				VTKAQKKDGKKRKRSRKESYS
İ						VYVYKVLKQVH\PDTGISSKAM
						GIMNSF\VNDIFE\RIAGEASRLA
						HYNKRSTITSREIQTAVRLLLPG
						\ELCKHAV\SEGTNA\VTKYTSS
23475	53843	Α	23610	93	421	
23476	53844	A	23611	2	1173	DHAAPRRRSRASVATVTRAGR
						CPTMSVRVARVAWVRGLGASY
						RRGASSFPVPPPGAQGVAELLR
						DATGAEEEAPWAATERRMPGQ
				1		CSVLLFPGQGSQVVGMGRGLL
						NYPRVRELYAAARRVLGYDLL
						ELSLHGPQETLDRTSQVQSPGT
						GIPRRVRAHVLHLSPQVIENC/V
						VS/AGLAFSPSFCALIFPNALL/F
			<u> </u>			WPGLYAVKIRAEAMQEASEAV
						PSGMLSVLGQPQSKFNFACLEA
		ļ				REHCKSLGPVNGQICKGGFLGH
						CSAETSFLPQALRFLQKNSSKF
		İ				HFRRTRMLPVSGAFHTRLMEP
						AVEPLTQALKAVDIKKPLVSVY
						SNVHAHRYRHPGHIHKLLAQQ
						LVSPVKWEQTMHAIYERKKGR
						GFPQTFEVGPGRQLGAILKSCN
23477	53845	Α	23612	187	425	NAME OF THE PROPERTY OF THE PR
23478	53846	Α	23613	1	1258	MKAYLLLGAAPPLPWFPGQAV
		ļ		Ì		FRPCPPPAGLWRTAQLVELEHG
						PVPREDLIPCSRPFSISFLGPERV
						DGLAFPRIQALGIAAARGMEPK
						GAPGKSDSGNSRSAGGRVLGC
				1		QGAALEVECSAGRGGPVRLAV
						GCVLLRAPERGWRRGEGAARG
						IPAGRSSESLGALSPQVGVRSDI
						GINGNIWHPGVQTDFSSLKRSQ
						GLYAVKIRAEAMQEASEAVPS
						GMLSVLGQPQSKFNFACLEARE
						HCKSLGIENPVCEVSNYLFPDC
						RVISGHQEALRFLQKNSSKFHF
1						RRTRMLPVSGAFHTRLMEPAV\
	1					EPLTQALKAVDIKKPLVSVYSN
	1					VHAHRIQGIPGHIHKLL\AQQLV
						SPVKWEQTMHAIYERKKGRGF
l						PQTFEVGPGRQLGAILKSCNMQ
L	<u> </u>	$oldsymbol{ol}}}}}}}}}}}}}}}}}$	<u> </u>	<u> </u>	<u> </u>	AWKSYSAVDVLQTLEHVDLDP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23479	53847	Α	23614	2	1050	GRCPTMSVRVARVAWVRGLG
						ASYRRGASSFPVPPPGAQGVAE
						LLRDATGAEEEAPWAATERRM
	İ					PGQCSVLLFPGQGSQVVGMGR
						GLLNYPRVRELYAAARRVLGY
}						DLLELSLHGPQETLDRTVHCQP
						AIFVASLAAVEKLHHLQPSVIE
ŀ				1		NCVAAAGFSVGEFAALVFAGA
						MEFAEG\LRFLQKNSSKFHFRR
						T\RMLPVSGAFHTRLMEPAVEP
						LERKLLKAVDIKKPLVSVYSQR
				:		PTRIRLQAFPGNIQQSCLAQQLG
			,			LPK*KWEQ\TMHAIYERKKG/R
						RGFPQTFEVGP\GRQLG\AILKS
						CNMQAWKSYS\AVDVLQTLEH
						V\DLDPQEPPKMTGRGLKCNDP
	<u> </u>]		LCPPEERL
23480	53848	Α	23615	600	905	QAPASGSPWLSGGPQQVEDAG
						AGYGFAPGQPPPPPRTQPRSAC
						SRRAAG/TPVPRAPPAPSAG\PK
1						ARAPRSLGCGSLGRFSTGVRPD
]	L				KCIFPPETENAAGCF
23481	53849	A	23616	390	722	QAPASGSPWLSGGPQQVEDAG
						AGYGFAPGQPPPPPRTQPRSAC
						SRRAAG/TPVPRAPPAPSAG\PR
						AAAPRSLGCGSLGRFSTGVRPD
]		KCIFPPETEMSYLLFRNTVSRCR
						C
23482	53850	Α	23617	1	673	MGQVKGHKLCLDFTLLSPDGG
ļ						AAEGRCLELGWQAKEVEETIE
						GMLLRLEEFCSLADLIRSDTSQI
						LEENIPVLKA\KLTK\MRGIYAK
	}					VDRIKAFVKMVGHHVAFLEAN
						VLQAERDHGAFPQALRRWLGS
						AGLPSFRNDPTAPLRQVFEASP
						DPRLARTVTIQTQQPVRVSGPL
						SGQRKEKSLQKGAQVFQTNVP
						ETLLEGAAISPAPACGGPLTSGD
						GFRHVRK
23483	53851	Α	23618	1	651	
23484	53852	Α	23619	195	497	DQFLENTKKCDGLNQLSFDPA
						NNNKPLEFGRARGPLVTEGLIE
						NGGESSKKRRRTNSSSADNSRT
						LNVDSTAMTLPMSDPTAWATA
						M\NNFGMAPVRIVGH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23485	53853	Α	23620	241	1823	NDDRAKFTVEDHTMQRDFRW
						LWVYEIGYAADNSRTLNVDST
						AMTLPMSDPTAWATAMNNLG
						MAPLGIAGQPILPDFDPALGMM
						TGIPPITPMMPGLGIVPPPIPPDM
						PVVKEIIHCKSCTLFPPNPNLPPP
						ATRERPPGCKTVFVGGLPENGT
						EQIIVEVFEQCGEIIAIRKSKKNF
						CHIRFAEEYMVDKALYLSGNR
	İ	l				ELKKFLNKEDNYFRKINLTALY
						KTVLEERCEKRRGGYRIRLGSS
						TDKKDTGRLHVDFAQARDDLY
						EWECKORMLAREERHRRRMEE
						ERLRPPSPPPVVHYSDHECSIVA
						EKLKDDSKFSEAVQTLLTWIER
						GEVNRRSANNFYSMIQSANSH
						VRRLVNEKAAHEKDMEEAKEK
						FKQALSGILIQFEQIVAVYHSAS
						KQKAWDHFTKAQRKNISVWC
						KQAEEIRNIHNDE\LMGIRREEE
						MEMSDDEIEEMTETKETEESEL
						KKLEGIVDGSVEVSAEALRKLL
						NEYERQVKEEQKVGYRIWGTI
j						HNKGEGKKEEFAEELEKSSSER
23486	53854	Α	23621	996	1512	RPVHLASPAKEQQGTRKGRRA
						PARLPVPTTGRGLEKGPESQAP
						PSFRQATGRFSRSGSSGGPTPSK
			i			NSRDCSHTPQLPTPRGNPYVFPS
						PGHPEPSRPVEMPAPG\EPRTRQ
						LCRERTRKACPPKPRPPLGLPG
						DPTGPVTHHAPPVSPTGASGQE
						RRAEPGAVSYAHASATK
23487	53855	Α	23622	293	1140	PRGILHQDKNLVVINK\PYGLPV
						HGWPWG\QLCITDVLPILAKML
						HGHKAEPLHLCHRLDKETTGV
						MVLAWDKDMAHQVQELFRTR
						QVVKKYWAITVHVPMPSAGVV
						DIPIVEKEAQGQQQHHKMTLSP
						SYRMDDGKMVKVRRSRNAQV
						AVTQYQVLSSTLSSALVELQPIT
						GIKHQLRVHLSFGLDCPILGDH
						KYSDWNRLAPQKLSVGTLKKL
						GLEQSKARYIPLHLHARQLILPA
						LGSGKEELNLVCKLPRFFVHSL
						HRLRLEMPNEDQNENNEAKCL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23488	53856	Α	23623	1	1328	MAVDLGSHTPGSSGVTAATFR
23 100			25025		1.520	GRRLVNTREESVVLGAGGEVL
		l				SGVPGTWDALGPGASFGIKGM
						FPKGTVIMSTSPMCEGPVPRRK
						PCASSVGQQNSRSRKKAGRKTP
						TLAGRHVPVKIKKEAIQQKPLP
		1				ASSFSPALTRSGAIFGPCPALTS
		ŀ				APVLEYPLQKVVENEAFKVFN
						NRDEESKRQKQAEFRMLTSAIR
						GPAGPRGHSSTRKPPSNPPPPGA
						CFKCGNEGHRSRQCPNPGTTSQ
						EAELIALTRALTLAAGQQINIYS
		İ				NSHYVFHIVHSHSSIWKERGILT
						AKNTPAINGSLINKLLKAARLP
						QKVAIIHCRGHQTPDNPILAGN
						VLADQVAKQVALQPVQGQFLS
	1	l		1		LSLFSPLYSSEEKEDFRAQNLQ
İ						KQGPW/LQSQGARLLVPAACAS
		1				KVAEAGRKPAGRHVPLKIQKE
						AIWSSPFSVPMATRAAASTSPT
23489	53857	A	23624	475	622	IHSKVLFSASSSSC*NTNNSWFR
		1				ASIKKNPEKQNKNMQSAIPQHG
		1				LWKI
23490	53858	Α	23625	17	992	GRGEVLRGRGRGQLDAVLSRE
						HCGGSRQSEQPWSRRSSSSSSR
						RRLRSAAEPAMALSMPLNGLK
						EEDKEPLIELFVKAGSDGESIGN
İ		ŀ				CPFSQG\LFMILWLKGV\VFSVT
						TVALKRKPADCQNLVPGTPPP\
		l				LILFSSEVKTDVNKIEEFSEEVL\
						CPPKYLKLSPK\HPRSNTAGMDI
		ŀ				FA\KFSAY\IKNSRPEANEALER
						GLLKTLQKLDEY\LNS\PLP\DEI
			1			DENSMEDIKFSTRKFLDGNEMT
						LADCNLLPKLHIVKVVAKKYR
						NFDIPKEMTGIWRYLTNAYSRD
						EFTNTCPSDKEVEI\AYSDVAKR
						LHQVKSRLLKEVSFMSSP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
23491	53859	A	23626	1	1128	MALASGEGLPVALSHGRRGNG QERVPYLKLRDTCAYRVAWRC
						NRLARAKVKRTSYRSGYGNGF
						EPGVEPDCVVLLALGVENDLV
İ						RISSKAEKYSLYEFSFPLRVSRE
						CEEECRKEPWFNPGQKPSVLIT
						SGQILGQEEQKTLLFPGLLSLVF
						QFNVENSDLRCLSLAENLEISAL
						LSLSCSSVIRLILINLEVTISPPITI
						LKHAPPLTSLEASSSSSGIEKKIH
1						NNFRQGPFPSSATRSTFLMNHT
						SFDSYVVVSNIVPRLNLYPQVL
						DVLQSLHKIHQLASNAKWTIQG
	1		•			PAKPDLGLELGLHRYPDWEKE
			Ì			ERKKKRKRSVSPEKSITTHNKP
						TSSGIFY/RPRRCGCDLGRPPAP
						GPHPSSPPHSPPQPSRCPPV
23492	53860	Α	23627	3	453	SLGKDIRPRSARAACKGVGLWS
						GCFGKMAGSGVRQVTSTASTF
						VKPIFSRDMNEAKRRVRELYRA
						WYREVPNTGER*RLVRPAVLK
22.402	52061	_	22.620		695	HSYVANRSF
23493	53861	Α	23628	3	095	PYASRRKFRVDPRVRAWSGCF
		1				DKMAGSGVRQATS\TASTFVKA HFSVGDMNE\AKREGARVLTAP
						WYR\EVPNT\VRVHQFQLDI\TV
						KMGRDKVREMFMKNAHV\TDP
						RVVDL\LVIKGK\IE\LEETIKVL
		ł				KQ\RTHVMRFFH\ETEAPRPKDF
						LSKF\YV\GHDPLKSFSGKDAR
						WILFLGAQIKLTIQWSLCDRIPL
						MNQFSAASFCLSKWGPWVLHI
		ŀ				TCSFPFSWKCGVCCKLD
23494	53862	A	23629	82	441	SLPINPVPTACTLQPAGLLRPPG
23.17.	33002	`			' ' '	SDVRTPMLPPSAPSQPSYPTSRP
						PLWPTWPAGPVFRRSQPLAQPN
						LPRSYHQQAPGGQTPVALGLE\
						CPKATNQPFGTLQRDPKSPAKV
						PPFEEFSL
23495	53863	A	23630	3	504	DRSSSSTMSTGGDFGNPLRKFK
	· · · · ·	1				LVFLGEQSVGKTSLITRFMYDS
						FDNTYQATIGIDFLSKTMYLED
						R/TNVNSFQQTTKWIDDVRTER
						GSDVIIMLVGNKTDLADKRQVS
						IEEGERKAKELNVMFIETS/AKA
						GYNVKQLFRRVAAALPGIMPS
						VLFQIIPSQALTRYLC
L	l	Ь		L		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23496	53864	Α	23631	416	1239	GGGRSSSSTMSTGGDFGNPLRK
						FKLVFLGEQSS\GKTSLITRFMY
		l				DSFDNTYQATIGIDFLSKTMYL
		ļ				EDRTIRLQLWDTAGQERFRSLIP
						SYI\RDSAAAVVVYDITNVNSFQ
		İ				ANYKSGIDDVRTERG\SDVI\IM
						LVGNKTDLADKRQVSIEEGERK
						AK\ELNV\ILMETSAKAG\YNVK
				-		QLLSTCKQPALPGMESTQDRSR
						EDMIDIKLEKPQE\QPVSEGGLF
						LTNLPCHLQPFFQKLTGFWPPY
						SFIDCSVNIGLMPFPFSNNVLRFI
				İ		IAACPAWR
23497	53865	A	23632	1	493	PKPATAPPPPSL/RARFLCPQRRP
		1				ERAR/PRARGASPRPEKTAGAR
			1			KRRRGSCERPHPGG/RPPAPG*I
						SSRPTLGFFWILNFFFLKKLGRI
						KDVPWQDSTKEHLSVCRGLRG
						TWSVPSASDPLHPAWRRTCFRV
						PPTTGAREWWLHLPPPALQRPP
						DSRQCHRIPVC
23498	53866	Α	23633	1	560	MESIWAGLWTAQDTTVLADDF
			1			EFILWTLKRRPERAR/PRARGAS
			Ì			PRPEKTAGARKRRRGSCERPHP
l			ļ			GGLRRLPGARDQRAPPRSDAG
1						RSPGRLRRLPALLCAGPLSPGRE
				-		PRRWEPRAGVARPAGQQWGFF
İ						WPGNQQGASICSLILWFLLECG
			 .			SRSTGGPCSGTELGHFSRERPTV
				1		LLSRLGTVKRP
23499	53867	A	23634	32	327	LTQRLFGGLFTRTHMKFGALTR
1	ı					IG\DLPWE\INPLSS\CSLLPEKDP
ł						PMTSGPQTDQPKEHLTNFKSAA
						RRSSRTSSHRNLLHVLEIWQLD
						QGMPAARDF
23500	53868	A	23635	2	424	LTPFSDSAHLHPGEINSPVAHTK
		1				PVGGLFTRMRVTFGAKTQDRS
						TP\QETSPLSSPS\PVTSGPQTNQ
						PKEHLTNFKLGKQSFHSSHASL
						ATLOSSSLATLOSPCPSNSNSFS
1		1				SLVETRRRPGAVARLQSQLSGR
		1				LRGGRIA
23501	53869	A	23636	31	246	TALLLTQSLFGGLFTRTHMKFG
						AVTRIG\DLPWGINPLSSCSLLR
	1	1				EKDPLTISGPQTHQPKEHLTNF
						KSGPH
		Ш	<u> </u>		<u> </u>	1100111

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23502	53870	A	23637	125	464	TAVFLTQSLFGGLFTRTRMKFG TMTRI/RGDLPWEINPLSSCSLL HEKDPPTTSGPQTNQPKKHLTN FKSRKVERTKGLLKTHLTKLSH QLKKAWTILLPLSLLRIQACPR NAT
23503	53871	A	23638	1064	1304	TAMLLTQSLFGGLFTQMCMKF GAVTRIG\DLPWEINPLSSCSLL HEKDPPTTSGPQTDRPKEHLINF KSGACYTCRKSGH
23504	53872	A	23639	539	747	TAMLLTQSLFGGLLTRTRMKF GALTRIG\DLPWEINPLSSCSLL HEKDPHTPSRKIHIRPQGNHESQ KGL
23505	53873	A	23640	709	950	TALLLTQSLFGGLFTRTHMKFG AVTQIG\DLPWENQSPVLLCSLL REKDPPTTSGPQTDQPKEHLTN FKSGKWSFHSSPAF
23506	53874	A	23641	132	1277	MGKKVLGIASQLLGAVRCEPD GRRVLPEDWPECLSLVTHGTW FTRQGDKNIQMADNSFSDGVPS DFVEAAKNASKTEKLTDQVMQ NPGVLAALQERLDNAPHTPSSY IETLPKAVKRRINALKQIQVRC AHIEAKFYEEVHDLERKYAAL YQPLFDERREFITGDVEPTSDM ESEWHRENEEEEKLAGDRKNK VVITEKAAATAEEPNPKGIPEF WFTIFRNVDMLSELVQEYDEPI LKYLQDIKFHFQPNDCFANSVL TKTCKMKSEPDKADPFSFAGPE IMDCDGCTIDWKKGKNVTVKT IKKKQKRKGPGTVRTITKQVSN ESFFNFFNPLKASGDGESLDKD SEF/ASDFEIGHFFRER\PQAVLY FAGEALKDDDNFDEGEEEEDEL EGDEEGEDEDEVEINPNKEPSQ
23508	53876	A	23643	1	515	RVRHRSVSVGSTHASAHAYGE GSLRIIPVTRSPLVTMNHIVQTF SPVNSGQP\PNYEMLKEEQEVA MLGAPHNPAPPTST\VIH\IRSET SV\PDHVV\WSLFNT\LFMNTCC LGFIAFAYS\VKSRDRKMVGDV TGAQAYASTAKCLNIWALILGI FMTI\LLIIIPVLVVQAQR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23509	53877	Α	23644	243	643	NHTVQT\FFSPVNSGQPPQRMR
						CIKEEHEVAVAGG\PHNPAPPTS
						T\VIHIRSETSVP\DHVVWSRVQ
						TPSSMNPC\C\LGFIAIRLTP*SLR
						DRKMVGDVT\GAQA\YAST\AK
						CLN\IWAHDSGASVMTILLIVIPS
23510	53878	Α	23645	3	1498	RCNFLRSSRIRVHPTPAASTMPP
		1				KFDPNEIKVVYL\RCTGGEVGA
						TSALAPKIGPLGLSPKKVGDDIA
						KAMGDWKGLRITVKLTIQNRQ
						AQIEVVPSASAMIIKALKEPPRD
					İ	RKKQKNIKHSGNITFDENGNIC
						STDMR/HRSLSRELSGTTKEILG
						TAQSVGCNVDGRHPHDIIDDIN
						SENDKQPRKSKMRTSFSPQIPG
						HEHFHYLLLLQHVHCQLALTL
ļ						RARKEGSSTHHQKLSTYYFKEG
	1					NSKEPKKIQRCRNTELRKEEGG
						GGSGKAPGSEIKLESPACSGASL
						GEMTAWTMGARGLDKRGSFF
						KGPLPNTCGHFWEMVWEQKSR
						GVVMLNRVMEKGSVVEFVTEL
		1				FFYVALPKMFPVHYIVILLASKL
						EEVTFNFPPPPDLCTPFLQNYLR
						VAFQEVNSGCTGKTLLVRPYIT
						TEDVCQICVEKFKVGDPEEYSL
						FLFVDETWQQLAEDTYPQKIKA
		1				ELHSRPQPHIFHFVYKRIKNDPY
						GIIFQNGEEDLTTS
23511	53879	Α	23646	361	918	GCTGSRGTVPLSALGPQRIGPP
						GSCLQIKSWVNDICQRQPG\DW
						KGP*GFTVKLTHSRTGQAPDCT
						RVP/VLPSCP*FIQSPSREPPKRPE
						KKPEKTFKHQLGIFTFLNEICST
			1			LLRQMR\HRSLSPEKLSWNPT*
						RDSWGIAQFSGAVNVDG\RH\P
						HDI\IDDIQQWVLWKCPRQLST
						KENIFNKGII
23512	53880	Α	23647	1	642	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23513	53881	A	23648	894	2438	VETKRIQPQPPDEDGDHSDKED EQPQVVVLKKGDLSVEEVMKI KAEIKAAKADEEPTPADGRIIY\ RKPV\KHPSDEKYSGLTASSKK KKPNEDEVYQDSVKKNSQKLII NSSLLSFDNEDENDTEKSMENL TGKIRKLQLRIVDPTFLPKLSNK GKDPVAVARLETGKGNKGAFR VLRDISRWATLCTSPQHSSSEA KNIIPYQATQFQNEVHKWYKK KGGLKISKRKALFFLTQDIQKK LQKLALGPNTPTPDILKIASLVF YNQDQEEKERAQQNERQKEKL YWLSYKSASPFQVALRILSQET THKDELPTDLPNNPEVQAPGIP GNTIMAMPVAIQLKDPSSYPCR IIKVPEIASWIHRTQVKLWKSPE TSELEPTPSVIEYTREALEDLKF LFKQKDNSSAKTPGRAYAASPR DWTSIEAELHIFYQWDPNLKG MFKPASLLAKVKQDFPDIHKES PIFRAIFSNITLMGIAPICVTGKTI KKRTRQHHTTPKSNLREQASQ
23514	53882	A	23649	200	1727	GSLNWEGN RTATSLDSVLYKSDYMTWKYR VSFFYILSCRLLKK/QCTDNEHV YMELSKKLYKYCPKEWKKEAS KVRQYEVTWVVSKRGKDYILK HIPNMHKDQFALTASEAHLKYI KEAVRLDDVAVHYYRLYKDK REIEASLTLGLTMRGIQIFQNLD EEKQLLYDFPWTNVGKLVFVG KKFEILPDGLPSARKLIYYTGCP MRSRHLLQLLSNSHRLYMNLQ PVLRHIRKLEENEEKKQYRESYI SDNLDLDMDQLEKRSRASGSS AGSMKHKRLSRHSTASHSSSHT SGIEADTKPRDTGPEDSYSSSAI HRKLKTCSSMTSHGSSHTSGVE SGGKDRLEEDLQDDEIEMLVD DPRDLEQMNEESLEVSPDMCIY ITEDMLMSRKLNGHSGLIVKEI GSSTSSSSETVVKLRGQSTDSLP QTICRKPKTSTDRHSLSLDDIRL YQKDFLRIAGLCQDTAQSYTFG CGHELDEEGLYCNSCLAQQCIN IQDAFPVKRTSKYFSLDLTHDE
23515	53883	A	23650	182	423	RLSIMASSLNEDPEGSRITYVKG DLFACPKT\DSLAHCISEDCRM GAGIAVLFKKKFGGVQELLNQ HYKEKGLRTSQLYG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22516	152004	<u> </u>	22651	1,	240	
23516	53884	A	23651	435	249 494	
23517 23518	53885	A	23652 23653	163	578	DALAM ANTI MECHEDEACHAE
23318	53886	A	23033	103	378	RVTWLKKHLNSGNSDSVCMAS SLNEDPEGSRITYVKGS\LFEGP
						KTDFLAPCISEDCRMGAGIT\VL
						FKKKF\GGVQELLNQQKKS\GE
						V\AVLKRRWGRLYIFTWITKRK
						RAFAQSPLYENLTEEFRGQLKS
						HLSEEWSH
23519	53887	A	23654	1	2238	
23520	53888	Α	23655	26	1941	RDAGTKWQRLGHLSCHCRKQA
	1	1				AREIINMHFQAFWLCLGLLFISI
		1				NAEFMDDDVETEDFEENSEEID
		l				VNESELSSEIIYKTPQPIGEVYFA
						ETFDSGRLAGWVLSKAK\KVD
						MDEEISIYDGRWEIEELKENQV
				ļ		PGDRGLVLKSRAKHHAISAVLA
						KPFIFADKPLIVQYEVNFQDGID
			ĺ			CGGAYIKLLADTDDLILENFYD
		ŀ				KTSYIIMFGPDKCGEDYKLHFIF
						RHKHPKTGVFEEKHAKPPDVD
						LKKFFTDRKTHLYTLVMNPDD
						TFEVLVDQTVVNKGSLLEDVV
						PPIKPPKEIEDPNDKKPEEWDER
						AKIPDPSAVKPEDWDESEPAQI
						EDSSVVKPAGWLDDEPKFIPDP
						NAEKPDDWNEDTDGEWEA\PQ
						ILNPACRIGCGEWKPPMIDNPK
						YKGVWIPPLVDNPNYQGIWSPR
						KIPNPDYFEDDH\PILLTSFSALG LELWSMTSDIYFDNFIICSEKEV
						ADHWAADGWRWKIMIANANK
						PGVLKQLMAAAEGHPWLWLIY
			1			LVTAGVPIALITSFCWPTKVKD\
						RHSDLEYIIPDICIPQTKGVLEQE
						EKVEKAALEKPMDLEEEKKQ\N
						DGEMLEKEEESEPEEKSEEEIEII
						EGQEESNQSNKSGSEDEMKEA
		İ				DESTGSGDGPIKSLRKRRVRKD
23521	53889	A	23656	297	398	
23522	53890	A	23657	286	430	TAGARCSSRTSSPGSCFNCQKS
						GHWAR\NARSPGFLLSRVPSVN
						WIHI
23523	53891	Α	23658	405	524	ARMSVACLPGLFFICLRGDFKQ
						NLTAYCWKCYQL*RSLGTHS
23524	53892	A	23659	1	657	
23525	53893	A	23660	22	334	ALKHFCLCSLIFSVTTMKFLAV
						LVLLGVSIFLVSAQNP\TTAAPA
						DTVSSLLVLLMMKPLDAETTA
						AATTATTAAPTTATTAASTTAR
						KDIPVLPKWVGDLPNR

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
23526	53894	A	23661	1	270	
23527	53895	A	23662	1	284	
23528	53896	A	23663	111	281	
23529	53897	Α	23664	229	363 586	PUBLICA PRODUCED VIVIORIA
23530	53898	A	23665	250	380	PVRHGAFFQDKSSDKKVQTKG KRGAKGKQA\EVAN\QETKEDL PAENGETKTEESP\ASDEAGEKE CQCLINNHIPCLISGPCLPSCTIQ RNIFINYFVNASFLVALETFLKK
23531	53899	A	23666	3	478	APAVGLSSSGGGRSGSGKAARF REGCRRAAARMQQAPGTRLPR RHPARAFPAATMPKRKVSSAE GAAKEEPKRRSARLST\KLPAK V\EAKPKKAAAKDKSSDKKV\Q TKGKRGAKGKQA\EVANQETK\ EDLP\AENGETERLEEESSLLIEA GEKEKPKS
23532	53900	Α	23667	1	252	
23533	53901	Α	23668	328	414	
23534	53902	Α	23669	1	3156	
23535	53903	Α	23670	1887	2036	
23536	53904	Α	23671	3	212	
23537	53905	Α	23672	22	539	YEGIVMKPLIIVNPADCIGC\RT CE\VACVVAHPSEQELNADVFL PRLKVQPIWTAVSAAR*CCPFSS ENAP\CVGA\CPVGA\LTMGEQV VQTNSARCIGCQSCVSACPFGM ITIQSLPGDTRQQIVKCDLCEQR EEGPACVESCPTQALQLLTERK LRRVRQQRIVVSGENPL
23538	53906	A	23673	135	302	PIHMKTRQPRDQFSADCVFLQS TCPVPC*RNRL*TPGGYWRTRR DYLRCDRSVHV
23539	53907		23674	175	794	LSTHSQPIADAADPPPHLHTGEL LCSAHQADFAVSAEPARPANN NAATTAPVHAVAITPPAYPDCF RTLPKSLSHQDSTQRLYGIARL LLRISTSGPPPSRLKVQRLDSISA PV/MCHQCENAPCVGACP/VGA LTMGEQVVQTNSARC/IGCQSC VSACPFGMITI/QSLPGDTRQQI VKCDLCGALRDCVAVLRQRRN SRVVLDACTRA
23540	53908	A	23675	3	264	LISLPFPPDSLAHRPPSSAPTEPQ GLFTVAAPTPGAPSPPATLAHL LPTPAMYSLLETELKNPVGTPT QAAGTGGPAAPG/GRKMALEN PKMHNSEISKRLGADWKLLTD AEKRPFIDEAKRLRAVHMKEYP DYKYRPRRKTKTL
∠JJ+1	22709	T.	123070	<u> 12</u>	207	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23542	53910	Α	23677	1	2377	MDDTFLNYKYKNKCGPKHIKK
		ŀ]	KEEEEEEDNNSNDSKNKDKTLT
						TGSSGSAKIFLMRGGRRQLLLP
						TLSLCLLSVQRSPGSHESGPPQS
						MNPRLSPQGKYLKNDFLDVIG
						MCDIACHLVVELLVGNNGNFL
						AHAFAHVEVVAQQNQRENAA
				•		RPNGGSVMIGQGASLRTCQSRV
						LRVARGGPSPNPGESEQGLFTV
						AAPAPGAPSPPATLAHLLPAPA
						MYSLLETELKNPVGTPTQAAGT
						GGPAAPGGAGKSSANAAGGAN
				•		SGGGSSGGASGGGGGTD\QDR
						VKRP\MNAFMVWSRGQRRKM
						APENPKMHNSEISKRLGADWK
						LLTDAEKRPFIDEAKRLRAVHM
						KEYPDYKYRPRRKTKTLLKKD
				•		KYSLPSGLLPPGAAAAAAAA
						AAAAAASSPVGVGQRLDTYTH
	ŀ					VNGWANGAYSLVQEQLGYAQ
				*		PPSMSSPPPPPALPPMHRYDMA
						GLQYSPMMPPGAQSYMNVAA
						AAAAASGYGGMAPSATAAAA
						AAYGQQPATAAAAAAAAAAA
						SLGPMGSVVKSEPSSPPPAIASH
						SQRACLGDLRDMISITGRENIFV
						VGDPAACLLPIEMEETFLPWPL
						GFEDRLHLEGPRPVQEIPSFLRK
						VPNCGQWLLQTPSVEPADPWE
		[•			NPQAFPPSLQQEQPCLSLIVGIF
						VEAFGGRGRWGSKGKKPKRYR
			!			TVEWMRIHQPSICCLQETHLTR
23543	53911	Α	23678	1	399	

3111

SEQ ID		Met	SEQ ID NO:	1	l	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23544	53912	Α	23679	3	1216	LGNKRAGLRRADGEAPSALPY
1	i	l				CYFLQKDAEAPWLSKPAYDSA
:	:					ECRHHAAEALRVAWCLEAASL
						SHRPGPRSGLSVSSLRPTRKMA
		l		İ		TTFLAHEKIWFDKFKYDDAERR
1		l				FYEQMNGPVAGASRQENGA\T
		İ				VILRDIARARENIHKSLAGSSGP\
						GAS\SGTSGDHGELVVRIASLEV
						ENQSLRGVVQELQQAISKLEAR
						LNVLGKELRLAHRGTAPQTQH
						VSPMR\QVEPP\AKKPATPAEDD
		ł				EDDDID\LVGRDNE\EEDKEAA
						QL\REER\LR\HYAEKKAQESLP
						LGGQVPPSLLDVK\PWDDESDM
						AQLEACVRSIQ/LGTGRSWGAS
						QMVAVGYG\IRKLQIQCVVEDD
						KVGTDLLEEEITKFEEARARSV
				İ		DIASFSTRSEALKCVYVAHVRE
						GPCTRSKDLRPAKKKKKK
23545	53913	Α	23680	1	2066	MAAALRAPTQVSAASSRPPLPS
			,			PPETEGRVKGFLQPGPHCPAQC
						LCSGDDGSNEDLEQKREVIFPR
		Į.				PHEAPVAEHRQQTVGLRELVAI
				}		PKRKEETTAIVQVFVAFEDVAI
						YFSQEEWELLDEMQRLLYRDV
						MLENFAVMASLGCWCGAVDE
						GTPSAESVSVEELSQGRTPKAD
		ŀ				TSTDKSHPCEICTPVLRDILQMI
						ELHASPCGQKLYLGGASRDFW
	'					MSSNLHQLQKLDNGEKLFKVD
		ŀ				GDQASFMMNCRFHVSGKPFTF
						GEVGRDFSATSGLLQHQVTPTI
ŀ						ERPHSRIRHLRVPTGRKPLKYT
		ŀ				ESRKSFREKSVFIQHQRADSGE
		ļ				RPYKCSECGKSFSQSSGFLRHR
						KAHGRTRTHECSECG\KSFSRK
						THLTQHQRVHTGERPYDCSEC
						GKSFRQVSVLIQHQRVHTGERP
						YECSECGKSFSHSTNLYRHRSA
						HTSTRPYECSECGKSFSHSTNLF
						RHWRVHTGVRPYECSECGKAF
				1		SCNIYLIHHQRFHTGERPYVCSE
						CGKSFGQKSVLIQHQRVHTGER
						PYECSECGKVFSQSSGLFRHRR
						AHTKTKPYECSECEKSFSCKTD
						LIRHQTVHTGERPYECSVCGKS
						FIRKTHLIRHQTVHTNERPYEC
						DECGKSYSQSSALLQHRRVHT
						GERPYECRECGKSFTRKNHLIQ
		<u>L_</u>				HKTVHTGERPYECSECGKSFSQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	1	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23546	53914	A	23681	1	468	MAAQVGAVRVVRAVAAHEEP
						DKEGKEKPHAGVSPRGVKRQR
						RSSSGGSQEKRGRPSQEPPLAPP
						HRRRSQPPTSWAAAVNDM/PAP
						TVPGPVEPLLLPPPPPPSLAPAG
						PADDAPLPAPSTSGLFTFSPLTA
						SAAGPKHKGTQGAAQAPSPPRP
						RW
23547	53915	Α	23682	950	1326	RPESQRANGVDSGPNLKTVPQP
						DTRKGSVLKWISKRGKPLAVEI
						EESHCL\CLPLRTECLGIKP\IVH
						LFSCTRPVIVPSLELNYDVDSIA
						HMFVADLLLMITLPSYDIPFYC
						LVFQNLLVLEFQYL
23548	53916	Α	23683	271	918	NQGLRDLGLCRICLVNKMFTSS
						ILGKGHRHSLVSIKQEHSALRK
						AAG\PLPLKVGYC\QGFSPCDSL
						KYG\SWDEKDLTVPQPDTHKGS
						VLRWISKRGKPLAVEIEEG\HCL
		}		-		PELPPGELECPG/ILKHGLYHWS
						SSEMGE\NRPMVG\ARHVYSNA
						ALLFFTALRIFGQRRNINLGLRA
						HPG\IVPPLELNYDIDSSAHM/FF
23549	53917	_	23684	1	942	LADLLLIITLPSYYIPFC
23550	53917	A	23685	346	1753	TWIPLSRTGRRFPVFSCSSSRYV
23330	33918	Α	23003	340	1733	AAISFHKRWKEGRCFGRTRGE
						MGWEGRSRSLKELNTRNDWNP
		1				GERELDFWRKITRFKEAASWTL
						EALEEAGSTARRRVWLEDGRR
						TCEILSVLCTSSLSSNCASLDCV
		1				TYKALSTTRLYDSVYWNKKPE
						RSSSPATEQSWTENDFDKLREE
						GFRRI\NYSELKEEARTHGKEVI
						NLEKK\LDERLTRITNAEKSLKD
		1				LMELKTMARELRDECTSFSSQF
		1				DPLEERVSVMEDQMNEMK\QE
						EKFREKRIKRN/EKQSLQEIWDY
						VKRPNLHLIGVPESDRENGTKL
						ENTLQEIIQENFPNLARQANIQI
						QEIQRTPQRYSSRGATPRHIIVR
						FTKVEMKEKMLRAAREKGRVT
		1				HKGKPIRLTADLLAETLQARRE
		1				WGPIFNILKEKNFQPRIAYPAKL
		1				SFISEGEIQYFADKQMLRDFVTT
		1				RPALKELLKEALNMERNNWYQ
		1				PLQKHAKW
23551	53919	Α	23686	73	327	GALQPTAALWEPLSGLAKAAA
	1	I .	1	1	1	1
						RSLSLQGGVEGDARAGTGAAR
						RSLSLQGGVEGDARAGTGAAR GACGPAGVPGGRGLGGPALGA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23552	53920	A	23687	322	402	DQEPTNSGHIGNLEQCLGHGYQ R*LFP
23553	53921	В	23688	545	3063	
23554	53922	A	23689	1	579	MKHEKRDRCAELTAILLNEQAS PMCTIVLGYTGTSTITLPRTSCH FAPRLSNSQQHSSYQVGHKPES YCVRNWWVLGLTDFKNEAAD PHNSGGQLASPSGSRTGAAGG\ VPASPAPCARTPQPLGGRWDW AP\GTGGGIRRGGSGCTGTHGG GGRLRHGGLQSGGLPRGKAAK ARREIESSAGYSNSSSKCSII
23555	53923	A	23690	399	592	KCIHFPGPTPKFVCGVIVVTAVP /TFGM*TVYV**LPFSILAQNVG *RDDCDPGWKLNLFYAKKKK
23556	53924	A	23691	3	268	DRV/RPLRKNFQRNDQTAAFTV HENPLLCSYRCWYPGKQEQKK IKPQRGGKNRAEKLETLKSRAP LLFQRNAVPHQQRNKAGWRM TLTS
23557	53925	A	23692		484	CSFFLVSMVFTFWHDFAAAGTS CSFPCLVLSFRSSFRAGLVVTKS LSICLSVKYFISSSLMKLSLAGY EIVG*KFFSLRMLNIGPHSLLAC RVSAERSAVSLMGFSLRVTRPF SLAALNIFSFISTLVNLTIMCLG VALLEKYLCGILCVS

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
23558	53926	A	23693	182	2076	MQEWLRLTARKTELSRDSRSC
23330	33720	^	23073	102	2070	GQLKTSLLISLLVESTQGRH*SF
1						L/VWTQTHSGGDGWRVAKAW
						1
		l	·			RPKYHDTQLGGEKEEPQGHHA
				1		SDCRWRAQPHCRTLSAAVVNV
ļ		1				AGFTVTFELRCLTLMKMREKD
				į		HHLCDNDIVLFPPDTAFPLDCL
						LSCIHSTVHGLLRWNAIQYLAS
				i		RSESHPERSEGGTPDRNGHPAL
		l				SLASFNGAQQDRRTVQSLGEKP
						GLPEEMAAFMAGTEKEQNELG
1						IEFFQLIIPASKRTHSENALFSHS
						GDDTECSRSGQGSPPSYSRGPP
		l				RRTFILATQPLCSKEAQTSPHEQ
						PTRTEKNRGSHLYHKSPCDHAS
1		1				PGMSVTMSGMQQALPRQLTTL
1		ļ		<u> </u>		TFYPPAFRSIKIFRIMILKTEMLH
		1				THARSQEALLKDSSKTMLHSTL
		Į .				PSCVRKSNYHHRERKIKTKEAF
ŀ		l				PAKIHRSRNQGMKVEVAPLTIT
	ľ	1				SSDPLAEFLLPVTVTLCSASLEV
		İ				LAPEGGMLSPGDATMIPLNWK
		1				LTLPPGHSGLLLPLSQQVKKRV
		l				PVLVGVIDLDYQDEIDYYSSMK
						ERSSSPATEQSWMENDFDELRE
]				EGFRRSNYSELREEIQTKGKEV
						ENFEKNLEECITRITNTEKCLKE
1		1				LMELKTKVRELREECRSLRSRC
	ľ	l				DQLEERVSAMEDEMNEMK
23559	53927	Α	23694	933	1118	RVERRRLQTIKLLRATGGNSNQ
	ŀ	l				RQRS*KL*KKFRRMYN*NNQY
		l		İ		REVLKGADGAESQGSRTT
23560	53928	Α	23695	2	250	WCHTKLDGLSLFGESVALEEA
		İ				GGQTALWLMRFLSRKANYVD
	ļ	İ				KITTREVGHRE*LMSSRLHSR*
	}					WAASSVCSSSCVLLPSSHS
23561	53929	A	23696	1017	1230	
23562	53930	В	23697	62	1069	
23563	53931	Α	23698	216	582	ILVPTLFWLGEFLLRDPLLV*W
		İ				ASI*G*PD/AFSLAALNIFSFISTL/
					:	RESDNYVSWSCSSRGVSL*RSL
ĺ		l				YFLNLNVGLPCQIGEVLLDNIL
			}			QSVFQLGSILPVTFGYTNQMQI
						WFHFIHFIFHH
23564	53932	A	23699	434	681	TLACLARLGKFSWIISCRVFSNL
						VPFSLSLSGTPIRRRFGLFT*SHI
						SWRLCSFLFILFSLNFPSRFISFV
						SSSITDTLSSS
23564	53932	A	23699	434	681	TLACLARLGKFSWIISCRVFS VPFSLSLSGTPIRRRFGLFT*S SWRLCSFLFILFSLNFPSRFISI

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23565	53933	A	23700	290	1231	RSFRAIGEVGSIRGWGGGYAAQ
						LQSPSRLPVLGGGKKGGSRGV
		-				GQGSSRAEVGLGSQRERSGTCG
		ŀ				AEAAAAAAQLIAGTGSSLPYQG
						PGQNQTAIAPTCAPLSPCKAAA
		1				VIMGNIFGNLLKSLIGKKEMRIL
						MVGLDAAGKTTILYKLKLGEIV
						TTIPTIGFNVETVEYKNISFTVW
						GCGVARDK\IRPL\WRH\YFQN\T
		1				QGFDILWSDSN\DR\ERVNEA\R
						EEL\MKNAGAETSLRDAVLPFL/
		İ				
						FGNKTDLRNAMNAAG\IQDKL
						G\LHSLRHRNWYI\QATCATRG
						N\GLYEGLHWLANSVPKQEVES
23566	53934		23701	164	208	RTALTKHPNRP
23567	53934	C C	23701	1	381	
23568	53936	C	23702	43	276	
23569	53937	C	23704	1	462	
23570	53938	C	23705	38	121	
23571	53939	A	23706	77	202	QKNYTQRPLPLHGLLELEVTCR
233,1			23 / 00			CWSWQPLLEPLELRP\RLSD
23572	53940	A	23707	123	209	
23573	53941	С	23708	1	411	
23574	53942	Α	23709	1400	1809	SSNPLSRPPLPTPVTPCPLLRLSL
						LRACCCCLFLPLRFFSCPLLCFL
						LCFFFSVCSPAHKNFKMPKPQW
						SGIPTGPPPSGSCFKCQKSGHW
						A\RNACSPGFLLSHVPSVWDPT
						GNQTVQLAWQPLLEPLEL*P\R
						LSD
23575	53943	В	23710	76	358	
23576	53944	В	23711	1	606	
23577	53945	С	23712	272	388	
23578	53946	Α	23713	1	259	PGVRGAQGGPSIPRQCEESAIGP
						KFAWIISSFHWKQGMFNMEAP
		Ī				SSLFFVNMCAVKKQTTWGRPD
						RGFIWQPLPEPLEL*P\RLSD
23579	53947	Α	23714	684	917	GDFGLRQWGFLNIQSCHLQTGT
						I*LPLFLIEYPLFLSPA*LPWPELP
	:		İ			TLCCIGVVREGISVLCQFSKGM
						LPVFAHSV
23580	53948	A	23715	618	683	
23581	53949	A	23716	306	710	DRVSLLTPRLGVQWPDPGSLQP
						RPPGFKRFYCLSLPSS*DYRHEP
					<u></u>	LLPAYFFPYRIIGWSRL
23582	53950	A	23717	2	88	

SEQ ID NO:	SEQ ID NO: of peptide sequence	•	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23583	53951	A	23718	972	1967	FPPGRGRGPGRPAGCSRGRARA CPKPVDHVKRPMNAFMVWSR AQRRKMAQENP\QMHNSEISKR LGAEWKLLTESEKRPF\IDEAKR LRAMHMKEHPDYKYRPRRKPK TLLKKDKFAFPVPYGLGGVAD AEHPALKAGAGLHAGAGGGLV PESLLANPEKAAAAAAAAA VFFPQSAAAAAAAAAAAA VFFPQSAAAAAAAAAAAAA VFFPQSAAAAAAAAAAAA VFPQSAAAAAAAAAAAA CYFPQSAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAAAAA VFPQSAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
23584	53952	A	23719	2	144	RPQRRP/LPPPTGSPLKHPGGAS GAAGAPIGEPQHPGAVVSVRA WSLCS
23585	53953	A	23720	219	897	PGRAGGTLGLGSRRDGTREKG VEETPHRLHARTDTTAPGCCGS PIGAPAAPEAPPGCFEG\PRWVE GVKKASPPLPAELIGWLVNFPL ARPVRPPSLSPGPPRRPPEGVCG SWPGRSEGERGKAKNTAESTPR QNLHRKYTIIVTHYQRNRKGQ NISFSLVCEFHPSLADWTTVTA LTWTSPKDNSLTEESWLREGN VHRGMDEYVRAQSFLILMTLE NSGGSVVSK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23586	53954	Α	23721	937	2730	ТИРНСИДНҮҮНИННИСИНСИ
ļ					ŀ	НННННСННННҮНННННСКОН
						НҮННННСННСННННННСНН
				·		ННҮНННҮНННСНОННҮНННН
						НСНОННҮНҮННННСНОННҮН
						ннннсноннунунннснонн
		ĺ				YННИННСИДНИУНИНИНСИД
						ннунннннсноннунунннс
						НННКҮНННННСННСННННН/Р
						ннининнсниннуниннис
						ннинуннинисниннунунс
		1				ннннүүннннсинсинниүн
						ннинсидинунинсинсин
			}			НҮННИННҮНДИНҮНИННСИ
						нсниннини\риннингисин
						НҮҮННИНИННИНҮНННЫ
		l				нгнсниналинининнин
		l				нннсннсүнннүнннүнсн
		1				ННҮНННННҮНРҮНҮНОНННН
						YHРННҮНННҮҮНСННННОҮН
		İ				НРИНСИННИНУНОННУУРІТІІ
						TTTITTITTIIITVTTTTFTTTTNT
		ŀ				TAPTTTPVTAITTTTTTTTTTTTTT
						TPITITTITNITTTTVTTTVIITTTF
						ТТТІНРҮНННННННРУТІТТТІТ
		l	-			TATIITTTTSITSITTITTTIVTTIII
						ТТТААТНННҮНРҮҮННҮННН
						ннннтнѕүүнүннннннсг
						нүннинүнин
23587	53955	Α	23722	2348	2571	
23588	53956	Α	23723	4919	5142	
23589	53957	Α	23724	2	370	
23590	53958	Α	23725	588	701	
23591	53959	A	23726	295	521	
23592	53960	A	23727	2837	3061	
23593	53961	A	23728	272	436	
23594	53962	Α	23729	239	514	
23595	53963	Α	23730	105	367	TGRPPELLLHLRSLSPPLL/CSSP/
						TPPPQLPPPPVSLASGYPLLHPL
						ARYPACPLPPRRARTEQRGGEQ
						PGVGASPLPIPGRWPGTEAT
23596	53964	Α	23731	999	1145	
23597	53965	Α	23732	1084	1893	

DHPENRIYRRKIEELS KTKGDGNCFYRAL\AI GKSREIFKFKERVLQT AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
DHPENRIYRRKIEELS KTKGDGNCFYRAL\AI GKSREIFKFKERVLQT AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY
KTKGDGNCFYRAL\AI GKSREIFKFKERVLQT AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY
KTKGDGNCFYRAL\AI GKSREIFKFKERVLQT AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY
GKSREIFKFKERVLQT AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY
AGFEEHKFRNFFNALR YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
YSVVELVEKDGSVSSL P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
P\GASDHIVQFLR/LCF RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
RADFFRHFIDEEMDIK VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
VEPMATECDHIQMTA ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
ALQVEYVD\GWIPALT AATPSVYMLYKTSQY DKH
AATPSVYMLYKTSQY DKH
OKH .
TML ADIZOUDEENCH A
TMLARKSIIPEEYVLA
RKPRIRDRLPKARFIA
LAHKNIREQGRFLQDI
LKWRHTLVIFTMSFLC
IMWWLVAFAHGDIYA
MEKSGLESTVCVTNV
LFSIEVQVTIGFGGRM
LAITVLILQNIVGLIINA
FMKTAQAHRRAETLIF
VRNGKLCFMFRVGD
SASVRIQVVKKTTTPE
IQLDIPVDNPIESNNIFL
HVIDKRSPLYDISATDL
VIVILEGVVETTGITTQ
EEIQWGHRFVSIVTEE
DYSKFGNTVKVAAPR
DEKPLPSLFRPSQKSEL
RKRNS\HRRNNFHRRN
N\SSLMVPKVQFMTPE
ES
VTSVCLTSAAASCRH
YLVS*KLPM/WSRRRK
S/V/CIVFWLASWMMS
DWVTIIAQQVLMNVV
CRMNLLEIFLRWLMR
RHFSQLLNVPITRPFS
ΓWVSARLMSYVLLMC
SCLTTPPHCSFSICFVI
KGSSLMYVFCLPNTAI
ORCRAATSRGSSRVDL
VGPSVPLVSSV**SDW
MTMMS

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23603	53971	A	23738	581	1434	KLPMGVAVVKLLHLPKVIIVFW LASWMMSRWLHYPDWVTIIAQ
						`
l						QVLMNVVSVMQADCRMNLLEI
						FLRWLMRRASDICSSIHGD*EPL
						L/YAVFGKQNTYIRLEPFKINVL
						EQITKHIEKLQCGGVVKQLSRR
						GNNQHISSTYDINRADTQVRRA
						VNNYDIIVMSNSFNGQSEHQV
						WIDRKAGECLLHEYEDLVPIRD
						TLRLFPGGRYLPRAKHVAPSEP
						DPEQDEQKLRFCKRHLYGKQP
						RSPVERVWDHGPTRIVGLIISLG
		<u> </u>				PRSHSYRRSDNQTGTTVPLVSS
23604	53972	Α	23739	515	774	LFPRLLSCLTTPPHCSFSICFVIC
						SRTLILKGSSLMYVFCLPNTAIS
						GTTVPLVSSV*LLVWDHGPTRI
						VGLIIRLGPRSHSYRRS
23605	53973	Α	23740	322	763	CRMC*LFPRLLSCLTTPPHCSFSI
						CFVICSRTLILKGSSLMYVFCLP
		l		1		NTAIVMALSPRGWRSKFGMPV
						EDTSLSTPAAPMVDSLIARVGV
		İ				MARGNAITLPVCGRDVKSTLE
						VLRGDSVEKTSRVWSGNERDQ
						ELLTEDALDDL
23606	53974	Α	23741	1	127	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEK*RRL
		l				SPLHSLMMTARKRGK*PGAGE
23607	53975	Α	23742	9	437	WLGLGNWKERLGTQTRMIWS
						QKTARRLCRTFSGIPAYSMCI/CI
						GMAERMWCDRNRERHTVSSSG
į		1		1		GNRLPNPDWAPGPPRIVGLNIS
						LGPTVPTRIVGLIISLGPRSHSYR
i		1				RSDNQTGTTVPTRIVGLILVLGP
						WSHSYRRSDY
23608	53976	Α	23743	766	767	LLTARLTWVSARLMSYVLLMC
						*LFPRLLSCLTTPPHCSFSICFVI
	•					CSRTLILKGSSLMYVFCLPNTAI
						GGSSRVDRCRAATSRGSSRVDL
						TL*KYSLVGPSVPLVSSV**SDW
						PLKLLLMTMMS
23609	53977	A	23744	573	1085	VCHQILTHLPALLSGARMMWD
25007	133711	l' `				QLWRGRP*RRCQQPQCGG*WH
						CFWATAHHSPLDSTLNHLHCLP
}						RGP
23610	53978	A	23745	2	379	WLTRSSCSWRLPPRRKKKGAL
23010	33910	^	23/43	-	313	V*PVWLPPGGVAVLLQLGCCV
						LQDVSCKRLCYNRKGQSEIVL
						MS*ITMWKLFPRPPLLPFSTCIC
						*KTCGKRGRSK*KVDILVLSIRF
1			L			SAIKLEPLDCHENAWL

SEQ ID	SEO ID NO	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23611	53979	A	23746	575	576	I TGIDSDQLNGRDWRRALSVSPA
						PDSGFPRCNGDEQAGI*LLGTG
						WAEKCHRLAFPPAGSTAGR
23612	53980	A	23747	11	723	MKLMETLNQCINAGHEMTKAI
-3012	33700	``	23/4/	1	7-20	AIAQFNDDSPEARKITRWRIG
						EAADLVGVSSQAIRDAEKAGR
		ŀ				LPHPDMEIRGRVEQRVGYTIEQI
		ŀ				NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKGGVYKTSVSVHLAQ
						DLALKGLRVCSWKERPPGNSL
		ŀ				NVSRMVWDHDPTRVVGLIIGL
		ł				GPRSHLYCRSDYQQRS\TSVCG
						CMPAVDCCCVLLIHNILRTVM
						WTKYLVTQAVPAR*PGCIRCKC
23613	53981	A	23748	1	1101	
23614	53982	A	23749	5	205	YCSSLSFTAAITSRAFLPDRRITT
				-		PPATSPSPFSSAMPRRISGPIWTF
						ATSRR*TGTPFSLAFKTM
23615	53983	A	23750	1	761	
23616	53984	В	23751	56	2188	
23617	53985	A	23752	627	1272	
23618	53986	A	23753	448	2218	VRRQRSDPERSDARMVRFCNL
						YI*RKNPFILH*LFR*TLRQTKPD
						NSAGKCVKI
23619	53987	A	23754	171	959	AASRGPDAPGLEMAGNCSWEA
						HPGNRNKMCPGLSEAPELYRR
						GFLTIEQIAMLPPPAVMNYIFLL
						LCLCGLVGNGLVLWFFGFSIKR
						NPFSIYFLHLASADVGYLFSKA
						LFSILNTGGFMGTFADYIRSVCR
						VLGLCMLLTG\VLLMPAGWDE
}						RGVRSERAWCTVETGRVHRSR
		ł				VRVSRNTARDPANGLGHLLQS
						LGMPALEPGPTLGSALRWEFLQ
						TASPQLPVQPLHPAAATPQRSL
	<u> </u>	 				RGRGRGPKARALGLGPRSTSYR
23620	53988	IA.	23755	664	838	
23621	53989	A	23756	2732	3007	OLIVE A CAMUCO CONTROL A POST
23622	53990	Α	23757	1522	1706	CHYFACMWSGCEVYS*SAPG**
						C*EDLSGLVRCRCLHSNAETPI*
02.600	52001	+	22750	1527	705	FAANADTSPRGPFLCR
23623	53991	Α	23758	537	795	ADCPLHMGRPCLID*GPCRTE*P
					1	RCAVVCLLWIAAVSCLSTT/SLR
					1	TGI/DGPKYLVTQAVPARYPGCI
22624	52002	 	22750	2074	2276	RCKCVAVDELASSDMRLPCS
23624	53992	A	23759 23760	2074	2276	ECWINCEL CAEL LILOD* A SCIL
23625	53993	Α	23/00	36	206	ECWIWCELGAELHHQP*ASSH WKSSYRQQSSLPRCRASQRTLH
						* *
23626	53994	A	23761	444	725	LLLQGWAVLGPLE
23626	53995	B	23762		1212	
23021	22773	լո	43704	1	1212	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23628	53996	A	23763	11	972	RFSKSLLDSGVQLASPSGSRTG
		ĺ				ATGGAAHSPARAPALHSPWAV
						DGTRGRLAGTAGLWSPRPAFH
						FKVSLSLPFSCVSSPFPGPPLGPS
						PRKHQMADDAGAGVGGRRPR
						GPRMG/NRGGFRRGFGSGIRHQ
						GCGCGQGQGQGRGVRGGKAE
						YKEWMTVTKLGHLVKDIKIKS
	Ì					LEEIYLFSFFKTSSSLKDEVLKI
						MLVQKQTHTSQCTRFKAFVAT
						GDYNGHISLGIKCSKEVATAIC
						GAIILAKLSIVPACRGYWGNKIS
						KPHTIPCKVAGCCGSVLVCLIS
						VPRGTGIVLAPCEGDRGLRKSS
						PASLSPESCKDGSHSRSF
23629	53997	A	23764	2	260	LLEECRSLRSRCDQLEERVSAM
25025	23777	 	25701	[200	EDEMNEMK*EGKFREKRISPKR
						APERSAKHGKEQPVPAAAKSC
						QNVNTIETRKKLHQLTSKITS
23630	53998	A	23765	3	1150	Q.,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
23631	53999	Α	23766	1	1521	
23632	54000	Α	23767	1	753	
23633	54001	Α	23768	1	3351	
23634	54002	Α	23769	1	1740	
23635	54003	Α	23770	1	1353	
23636	54004	Α	23771	1	1932	
23637	54005	Α	23772	1	1221	
23638	54006	Α	23773	1	1214	MAKKQNRKTGNSKKQRASPPP
						KECSSSPATEENWTENDFDELR
						EEGFRQSNYSELWEDIQTKGKE
						VENFEKTLEECITRITNTEKCLK
						ELMELKTKAQELRKECRSLRSR
						CDQQQERASVLEVLEVLAR\QL
						GRKRK*RVFN*EKRKSNCPCLQ
						TT*LYI*KTPLSQPKISLS**ATS
						AKSQDTKSMYKNHKHSYTPTT
						DKQRAKS*VNSHSQLLQRE*NT
						*ESNLQES*NWISSLHLIQKSIQD
						GLKT*TLDLKPSKP*KKT*ALPF
						RT*AWARTSCLKHQKQWQQRT
						KLTNWI*LN*RASA/PAKESTV/S
1						SEQATYNMGENFHNLLI*QRAN
						IQNLQ*TQTNLQEKNKQPHQKV
						GEGHQQTLLKRRHLCSQKTHE
1						KMSIITGHQRNANQNHNEIPSH
						TSYNGNH
	Ī					131NGNH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23639	54007	A	23774	1157	2085	QGRPTDFADFGTTIKQDFRLLG QTSVDRLLQLSQGQAVKGNQL LPVSLVKRKTTLAPNTQTASPR ALADSLIAAGTTGFPTGKRAVS ATQLMTQFSPAREQSWRKN/DF DELRKE/GFARSNYS\EKCLKEL MELKTKARELREECRSLRSRCD QLEERVSAMEDEMNEMKREG KFREKRIKRNEQSLQEIWDYVK RPNLRLIGVPESDAENGTKLEN TLQDIIQENFPNLARQANVQIQE IQRTSQRYSSRRATPRHIIVRFT KVEMKENMLRAAREKGWVTL KGKPIRLTADLVAETLQARRE WGPIFEEP
23640	54008	A	23775	1	1707	
23641	54009	A	23776	1	1899	
23642	54010	Α	23777	1	2781	
23643	54011	A	23778	1	831	
23644	54012	A	23779	1	3264	
23645	54013	Α	23780	1	882	
23646	54014	A	23781	1	1853	
23647	54015	A	23782	1	550	
23648	54016	A	23783	1	831	
23649	54017	В	23784	64	3436	
23650	54018	Α	23785	1	1422	
23651	54019	A	23786		773	MGKKQSRKTGNSKKQSATPPP KERSSSPATEQSWTENDFDELR EEGFRRSNYSELQEETQTKGRE VENFEKNLDECITRTTNTEKCL KELMELKAKAWELREECRSLR SRCNQLEERVSVMEDEMNEMK REGKFREKRIKRNEQSPQVIRD YVKRPNLRLIGVPESDGKNGTK LENTLQDIIQENFPNLARQANIQ IQEIPRTS/QRYSSEEHSIIGLIGR FPS*RTDMFRAPKKREQKRNNR KIQKEEGGRTAPTGI
23652	54020	A	23787	1	2775	
23653	54021	A	23788	1	2712	
23654	54022	A	23789	1	1590	MCVVONDVTCNEVVOCACDDD
23655	54023	A	23790		547	MGKKQNRKTGNSKKQSASPPP KECSSSPATEQSWTENDFDELR EEGFRRSNYSELREDIQTKGKE VENFEKNLEECITRITNTEKCLK ELMELKTKARELREECRSLSSR CNQLEERVPAMEDEMNEMKE D/GNRYDDRRQ/RKEEIMIKSW RFCCRHLIRVVRGWGPSKYLWI TMVVTLAGNPQ
23656	54024	Α	23791	1	710	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23657	54025	Α	23792	1	895	
23658	54026	Α	23793	1	1938	
23659	54027	Α	23794	1	1413	
23660	54028	A	23795	985	1219	LCSGSVERCTESYLHSSHCLVQ QAGVTGDEGGSCY*DPPSHHRS HPRKLKKLCEQEFSVTSAPWRK CRSLHGRFPGLP
23661	54029	A	23796	1	3570	
23662	54030	Α	23797	1	634	
23663	54031	A	23798	1	4551	
23664	54032	A	23799	1	4721	MHPSQRTPYQSRSPATIFATVST FKSHKSNLETKHPQQVPLSSLG LAANISSEDIESEAEDNGKWES VDRTSFFPALTSFQQSMIHKKI MTVFKSVEPEDGLGRPTLAFHE DRCQLSIQNSLDAGGGCPSMA APPAEVHELASGLISAAALLLSP AASGRVSKHQHIPTGFPDPPKA LHKGKLTTRKDIYTENPSVHHH HQRPKVDKTTKMGKKQNRKT GNSKTQSASPPPKECSSSPATEQ SWMENDFDELREE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, ∖=possible nucleotide insertion)
				sequence		
23665	54033	A	23800	1	2152	MDILGWASRAHKWKGRATGT
		l				HVFNLMREYLISLQCCASEQAA
1						SPEHGASFRVWKSLGFQKLILS
						LICPQCLQSLTRETWMRFKSKK
		l				TKGPDLRLVMDANKWEFRRH
		1	,			ALRPALTFPNEGGDLGVEALPV
		i				ALPPEAHVGGVVLPAALVRGA
		l				HGEQLPAIPGPEEFALPAVQAR
		1				AARPERQVVVIVVNEAGVDPA
		ĺ				GVEGLAPVPVLPGVRLEATAPK
		ĺ				QPPRLPARLRAPLPSGRKLNAK
		ł				CPRLRWGFFVQDGDGTELACL
		l				SEGDGGCCQEAPLDHRPRRLRG
						LAGSAGAREAREPRLRPGDWG
						NRIARVRAPEEAAAALSRPARV
						AAALAKTHRREHRARSQKLVM
						SSVQQGKLTTRKDIYTENPSVH
						HHHQRPKVDKTTKMGKKQNR
						KTGNSKKQSAF\PPPKECSSSPA
						MEQTWT\DNDFDELREEGFRGS
						NYSELREDIQTKGKEVENFEKD
						LEECITRITNTEKCLKERMELKT
						KARELREECRSLRSRCDQLEER
						VSAMEDEMNEMKREGKFREK
		1				RIKRNEQSLQEIWDYVKRPNLR
						LIGVPESDVENGTKLENTLQDII
						QENFPNLARQANVQIQEIERMP
						QRYSSRRATPRHIIVRFTKVEM
						KEKMLRAAREKGRVTLKGKPI
						RLTAVLSAETLQARREWGPIFN
						ILKEKNFQPRISYPAKLSFISEGE
						IKYFIEKQMLRDFVTTRPALKE
23666	54034	Α	23801	1	2109	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	I .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
:				sequence		
23667	54035	A	23802	1	2544	MSVHSWKPSWIRPIRSLSPKRPL
						GTQGTSEVPPDTHICQTKFHSLS
						SGPLLLSVHPVGKLSAVVGRVF
						IHILSQRERRRRRVISAFPSEVPG
į						SSHQGVPDSGCRSVGAHPVREP
						KQGEALPHSGSPREGKLTNRKD
						IHTKNPSVHHHHQRPKVDKTT
		ŀ				KIGKKQNRKTGNSKTQSASPPP
				ļ		KECSSSPATEQSWMENDFDELR
						EEGFRQSNYSELRKDIQTKGKE
						VENFEKNLEECKTRITNTEKCL
						KELMELKTKARELCEECRSLRS
						RCNQLEERVSVMEDEMNEMK
ļ						REGKFREK\RIKRNEQSLQEIWD
ŀ					1	YVKRPNLRLIGVPESDVENGTK
ŀ]				LENTLQDIIQENFPNLARQANV
l						QIQEIQRTPQRYSSRRATPRHIIV
İ						RFTKVEMKEKMLRAAREKDRS
						TRQKVNKDTQELNSALHQADL
			İ			IDIYRTLHPKSTEYTFFSAPHHT
						YSKIDHIVGSKALLSKCKRTEII
						TNYLSDHSAIKLELRIKNLTQSR
:						STTWKLNNLLLNDYWVHNEM
	:				:	KAEIKMFFETNENKDTTYQNL
						WDALKAVCRGKFIALNAHKGK
						QERSEIDTLTSQLKELEKQEQT
						HSKASRRQEITKIRAELKEIETQ
						KTLQKINESRSWFFERINKIDRP
						LARLIKKKREKNQIDTIKNDKG
					,	DITTDPTEIQTTIREYYKHLYAN
						KLENLEEMDTFLDTYTLPRLNQ
						EEVESLNRPITGSEIVAIINSLPT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23668	54036	A	23803	, , ,	1886	MKENFKHRHVVAFISSQEAQE DSREEEDRLESRFYFLFLLELSY KTTGLNREPTVSLNSKSGTPMD QNEHSHWGPHAKGQCASRSEL RIILVGKTGTGKSAAGNSILRKQ AFESKLGSQTLTKTCSKSQGSW GNREIVIIDTPDMFSWKDHCEA LYKEVQRCYLLSAPGPHVLLLV TQLGRYTSQDQQAAQRVKEIF GEDAMGHTIVLFTHKEDLNGG SLMDYMHDSDNKALSKLVAA CGGRICAFNNRAEGSNQDDQV KELMDCIEDLLMEKNGDHYTN GLYSLIQRSKCGPVGSDERVKE FKQSLIKYMETQRSYTALAEAN CLKGALIKTQLPKVDKTTKIGK KQSRKTGNSKKQSTSPPPKERS SSPAMEQSWTENDFDELREEGF RRSHYSELPEDIQTEGREVENFE KSLEECITRITHTEKCLKELMEL KTKARELREECRSLRSQCDQLE ERVSAMEDEMNEMKREGKFKE K\RIKRNEQSLQEIWDYVKRPN LRLIGVPESDAENGTKLENTLQ DRIQENFPNLARQANGFCRFRN HHQTGFSPAGANQRGPLAATLS GPGGEGQSAVARLTGEKKNHP
						GAQYANRLSPRVGRFINRSWH DRFPDWKAGTGLKSAIFAWE
23669	54037	Ι	23804	1	4134	Did i D Wild i G I G Did i i i i i i i i i i i i i i i i i i
		A		11		
23670	54038	Α	23805	1815	1849	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23671	54039	A	23806	1	4016	MGKKQNRKTGNS\KTQSAFP\P
						PKECSSSPATEQSWMENDFDKL
						REEGFRRSNYSELR\EDIQTKGK
						EVENFEK\NLEECIT\RITNTEKC
		1		1		LKELMELKTKGSRNYVKECR\S
		ŀ				LRSRCDQLEERVSAMEDEMNE
						MK*EGKFREKRIKRNEQSLQEI
			1			W\DYVK\RPNLRPIGVSESDGEN
						GTKLENTLQDIIPENFPNLARQA
						NVQIQEIQRTPQRYSSRRATPRH
						ILVRFTKVEMKEKMLRAAREK
						GQVTLKGKPIRLTADLSAETLQ
						ARREWGPIFNILKEKNFQPRISY
						PAKLSFISEGEIKYFIDKQMLRD
		İ				FVTTRPALKELLKESLNMERNN
						WTSKQIIQEDGYSEEEGWLCRA
						VIYSHTLQAIMAIVKATGNLQT
	1					DFANPFHTDDARQQFAMSYTV
						VGQGALPEDLSRCHLQTEAVQ
						RHPESLGCPLDLLHGMPVSSCF
						GYPQDTSSTV
23672	54040	Α	23807	2059	3153	
23673	54041	Α	23808	6	188	
23674	54042	Α	23809	3	1014	VACRPLPRRRAQPATFIPRVAV
						KMALVASVRVPARVLLRAGAR
						LPGAAFGRTERAAGGGDGA\RP
						\FGSQRVLVEPDAGAGVAVMK
		1		1	1	FKNPPVNSLSLEFLTELVIS\LEK
						LENDKSFRDVILTSD\RPGVFSA
						GLD\LTEMCG\RSPAHYAWVLE
						RPVQELWLRLYQSQL\VLVSAI
						NGA\CPAGG\CLVALTCDYTIL\
						AGNPRYCIGLNE\TQ\LGIIAPF\
						WLKDNPWRTPIGHRAAESA\LQ
				:		LGLPLPPPAEALQVGIVDQ\VVP
						\EEQVQSTALSAIAQWMAIPKP
						MAHK\LTKAIMRKGQGPPAWS
						RKGYADVQNF\VSFISKDSIQKS
1					1	LQMYLERLKEEKG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23675	54043	A	23810	1	2665	MTTSIRQFTSSSSIKGSSGLGGG
į						SSRTSCQLSGGLGAGSCRPGSA
						GGLGSALGGSSYSSCYSFGSGG
			:			GYGSSSFEGVDGLLVGGEKAT
				1		MONLNDRLASYLDKVRALEEA
				İ		DTELEVKIRDWYQRQAPGPAR
				İ		DYSQYYRIIEELQNKPPPSDDLT
						TLPSPLILTATVDNANILLQNDN
					ł	AHLAAADFRTKFETEQALCLSV
						EADINGPCRVLDELTLARADPE
				1		MHIENLKEELAYLKKNHEEMN
				ł		APRGQVGGEINVEMGAAPGVD
						LSRILNEMREQYEKMAEKNRK
						DAEDWFFSKTEELNREVATNSE
						LVQSGKSEISELRCTMQALEIEL
•						QSQLSMKASLEGNLAETENRY
					1	CMQLSQIQGLIGSVEERLAQLL
					}	CEMEQQNQEYKILLDMKMRLE
						LEITTYHRLLEGEDAQTVQHTP
		ļ.				GSPCVPEWAHDRCSLPAQLDE
						WSCQAQRGICGHRRENCWERIP
						SQSCFYRPTARSLTSLLGTMST
ł		1				CSHQFTSSRSMKGSCSIGGSIGG
						GSSRISSVLAGGSCHAPSTYGG
						ASCGLGGGYGGGFGSSSSFASG
				1		FGGGYGGGLGAGL
		l		İ		GGGFAGGDGLLVGSEKVTMQH
		l				LGDRLASYLDKVRALEEANAD
						LEVKIRDWYQRQRPSEIKDYSP
						YFKTIEDLRHKIIAATIENAQPIL
İ		l		İ		QIDNARLAADDFRTNYEHELAL
						QQTVEAGVNGLCREMLALRGQ
23676	54044	A	23811	222	548	TLPCGDHHHSQDPARLPSWHD
230,0		ľ.				LLQERSEKCGSQDWQRDSVQP
						GVSYQFCCVPWPAGRSPQPRH
					İ	CWGCCGTEASYDSGI*SLSTPG
ļ						GGQLQGSV*GPDGAGLQNSHR
23677	54045	Α	23812	2	165	
23678	54046	Α	23813	1	665	
23679	54047	В	23814	1	1673	
230/9	134047	l _D	123014	T .	110/3	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23680	54048	Α	23815	1	331	ARQLRTSAMTMPVNGAHKDA
						DLWSSHDKMLAQPLKDSDVEV
						YNIIKKESNRQRVGLELIASENF
						ASRAVLEALGSCLNNKYSEGYP
						GQRYYGGTEFIDELETLCQKRA
			;			LQAYKLDPQCWGVNVQPYSGS
		İ				PANFAVYTALVEPHGRIMGLDL
		l				PDGGHLTHGFMTDKKKISATSI
						FFESMPYKVNPDTGYINYDQLE
						ENARLFHPKLIIAGTSCYSRNLE
		İ				YARLRKIADENGAYLMADMA
		İ				HISGLVAAGVVPSPFEHCHVVT
						TTTHKTLRGCRAGMIFYRKGV
		l				KSVDPKTGKEILYNLESLINSAV
						FPGLQGGPHNHAIAGVAVALK
						QAMTLEFKVYQHQVVANCRA
						LSEALTELGYKIVTGGS/CVISG
						GAGKPG*FSHQDGTGTSQDGG
		ĺ				RRPFLHHTLSY
23681	54049	Α	23816	3	1709	SHPTPRETPTGVEGQATPSVLG
		ŀ				AAPASQAKATPTLGRPHASDR
						GEVAQLDEWSCQAQRGICGHR
		-				RENCWERIPSQSCFYRPTARSLT
						SLLGTMSTCSHQFTSSRSMKGS
						CSIGGSIGGSSRISSVLAGGSC
						HAPSTY/GGGLCVSFSRFSFGGA
						CGLGGGYGGGFGSSTNFWS\GF
						GGGYGAGLGAGFGGGLGAGL
						GGGFAGGDGLLVGSEKVTMQH
						LGDRLASYLDKVRALEEANAD
						LEVKIRDWYQRQRPSEIKDYSP
						YFKTIEDLRHKIIAATIENAQPIL
						QIDNARLAADDFRTKYE\HELA
						LRQTVEADVNGLRRVLDELTL
						ARTDLEMQIEGLKEELAYLRKN
						HEEEMLALRGQTGGDVNVEM
						DAAPGVDLSRILNEMRDQYEQ
						MAEKNRRDAETWFLSKTEELN
						KEVASNSELVQSSRSEVTELRR
						VLQGLEIELQSQLSMKASLENQ
						L\EETKGRYCMQLSQIQGLIGSV
						EE\QLAQLRCEMEQQSQEYQIL
						LDVKTRLEQKYCHPTAALLEG
						EDAHLSSQQASGQSYSSREVFT\
						SSSSSSRQTRPILKEQSSSSFSQ
23682	54050	A	23817	2	365	
	124020	<u>. </u>	12301,	<u> </u>	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23683	54051	Α	23818	1	464	RRWRICSTWTRSFATAWLPPAP
						SANFPRTDRSQGDRGAPAGFAL
	•					APILEFFLWDTSSQKFLQKPHCF
						MHGRERAKMGRRAQQESAQA
						ENHLNGKNSSLTLTGETSSAKL
	1					PRCR/QGGWAGDSVKASKFRR
į		ł	ļ			KASEEIEDFRLRPQSLNGSDYG
						GDIP
23684	54052	Α	23819	384	805	
23685	54053	Α	23820	3	392	MKYSAIQTLDGEIDLKLLTKVL
		l				APEHEVR*VQWWQQFPGLSTL
						AIPINPAELPGLALCLP*RPWNG
						REKVSLQLREKALSLSYPAKPF
		1				L\EDDVGWDWDHLFTEVSSEV
ŀ		1				LTE\WDPLQTEKEDPAGQARHT
23686	54054	A	23821	63	785	GPIRGPWGPGFGGWGGATPPAP
		1				TSLRRLRAAGLAPRRRAKMGR
		1				RAQQESAQAENHLNGKNSSLT
		l				LTGETSSAKLPRCRQGGWAGD
		1				SVKASKFRRKASEEIE\DFR\LRP
		1				QSLNGS\DYGGDIPI\IPDLEEVQ
						EEDFVLQV\AAPPSI\QIKRVMT
		ł				YRDL\DNDLM\KYSAIQTLDGE
		ł				D/IDLKLLTQK/VLAPEHEVRER
		1				NPSWQDDVGLGTWGPICSTEV
						SLRRSFTEWDP\LQTEKEDPGG
						Q\ARHT
23687	54055	A	23822	1	441	MSFNCSTRNCSSRPIGGRCIVPV
				•		AQVTTTSTTDADCLGGICLPSSF
						QTGSWLLDHCQETC/CGCQPLG
	İ	l				GISSVCQPVGGISTVCQPVGGV
		1				STVCQPACGVSRTYQQSCVSSC
		1				RR\PAKCVGASERIKTPRPASCF
		<u>L</u>				QDLPACCLSLNSSS
23688	54056	A	23823	1	538	PTRPLQQATCSPLSDIMSFNCST
		l				RNCSSRPIGGRCIVPVAQVTTTS
						TTDADCLGGICLPSSFQTGSWL
						LDHCQETCCEPTACQPTCYRRT
	1					SCVSNPCQVTCSRQTTCISNPCS
						T\PTAGRSP/CVSSGSQPLGGISS
						VCQPVGGISTVCQPVGGVSTVC
<u> </u>	<u> </u>			<u> </u>		QPACGVSRTYQQSCVSSCRRTC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23689	54057	A	23824	832	1885	AEWDAADKAHVGPGRSPROPL
		ĺ				PHVGRRRAAHTTGRGRRTAGC
		į	:		İ	APAPARTLCILSILGPWCVLPHL
		1				PTLCLGQAQAASCSQGAHGCF
						AGCSCT/GTSASHPV/CCKCPLH
						PVCTLADVHGCITSGFSVISLPQ
						TCAPWTSWRKWSLSDSWQVD
		ĺ				ACPESCEPPCCATSCCAPAPCL
		ŀ				TLVCTPVSCVSSPCCQAACEPSP
						CQSGCTSSCTPSCCQQSSCQPA
						CCTSSPCQQACCVPVCCKPVCC
		1				, -
					}	VPVCCKPVCCKPICCVPVCSGA
						SSSCCQQSSRQPACCTTSCCRPS
						SSVSLLCRPVCRSTCCVPIPSCC
				1		APASTCQPSCCRPASCVSLLCRP
22.500	-10.50	<u> </u>	22027	ļ. ——	2007	TCSRLSSACCGLSSGQKSSC
23690	54058	Α	23825	1	3097	MDEKRTRLQPEEADVPAYPKV
1						ATVPQSEKKPSKYPLCSADAAV
						STTPPGSPPPPPPLPEPPVLKVLS
						SLKPAAPSPATAATTAAAASTA
						ASSTASSASKTASPLEHILQTLF
						GKKKSFDPSAREPPGSTAGLPQ
						EPKTTAEDGVPAPPLLDPIVQQF
		l				GQFSKDKALEEEEDDRPYDPEE
						EYDPERAFDTQLVERGRRHEVE
						RAPEAAAAEREEVAYDPEDETI
l						LEEAKVTVDDLPNRMCADVRR
						NSVERPAEPVA
23691	54059	Α	23826	67	442	VKFFCLYLQSNEKPMKNFKQG
						SNNQLLMRSFWLLCGIDYKKPP
						PPPAPGPESVAPALSGPPPPAHS
						RGPRAVVRA\PPPPGPRHRSPAE
		1		}		PPPPPGTEPSVATYSIKASKRVS
						QQDRCHNLMKSQE
23692	54060	A	23827	3	657	GERSRRWGPACGPGAAEKTPP
						CDAGAAGAAPGAPGHAGNPGL
						GSGPDYD\GEATCMHLHPDDPC
į						PAGAGCCPGSFEVAFPDAAEK
						MKKSLHSLKKGCYSRVCDLPL
						DCPGCDSDSGRQAMFSCIVNFQ
						LPKEESPIPGSSQEESSGPSHSEC
						DSVGVDWRGQRVSRKLQAPNH
						SLVLLVTSLFQETTKSHLNRTN
						DAVSPRKFKRLGALYWGKVEA
23693	54061	A	23828	141	246	DA VSFRRI KRLUAL I WURVEA
23694	54062	A	23829	1	609	
23695	54063	A	23830	1619	1752	
23093	34003	<u> </u>	23030	1019	11/32	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23696	54064	A	23831	54	470	CSTMNPSEMQRIAPPRRQRHRS RAPSAHKMNRMVMSEEQMKL PSTKKAEPPTWAQLKKLTQLA KKK\LENTKVTQTPENMLLAAL KTVSTVSAGVPSSSEESDHRER AMMTTVVLSKRRGKCGEKKEI
23697	54065	A	23832	213	572	SDCYCVYVERS IQWYPYQDTIDFLHRIRKNFFKF HMEPKKSLHCQDNPKPKEQSW MHHTT*LQTMLQGYSNQKRMP PHLGSEERLCPAAPSGR*GAPLP GRCAIFQV*SDSLSAGVPNSSEE TATIENGP
23698	54066	Α	23833	296	583	
23699	54067	Α	23834	2452	4557	
23700	54068	Α	23835	3	266	
23701	54069	A	23836	31	499	ARVTAAVSAAAVAKRVWRDG FDCSGYYSLTTILLVQPTKRP\E GRTYADYES\VNECMEGVCKM YEEHLK\RMNP\NGPSI/THNTTS QLFDFID\DLADLSCLVYRA\DT QTY\QPYN\KDWIKDEDPTVLLS SGKAQQAREIIVVGSHWGGWG GAWDTGV
23702	54070	Α	23837	3	323	
23703	54071	Α	23838	366	892	PCVYSQFPAGEQCLKLHSPAEA SPPALEASEAQTRKAPECGSGV PEVAGGSCPCLLLCLPRHQAVP TQGPGTYSPCTSHQPHIFPRPAA PHHLGLLQNPHNAASCIQCLYP AGVATTMPRRKAEGDAEGD/K AKVKDEPQRRPAKLSAKPAPPN PEAKPKNAPGVTLSLRGTATRF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23704	54072	A	23839	201	1412	LWKSRLTFKLAMSRVPSPPPPA
	10/2	-				EMSSGP\VAESWCYTQIKVVKF\
				ļ		SYMWDPSITLAFCREEMG\EV\I
						KSSTFSSGANDKLKWC\LRVNP
						KG\LDEESKDYLVTFTCYLVSC\
		ļ				PKSEVRANFKFSILNAKGEETK
						AMESQR\AYRFVQGKDWGFKK
			•			FIRRDFLLDEANGL\LP\DDKLTL
						FCEVSVVQDSVNISGQNTMNM
						VKVPECRLADELRGLWENS\RF
		1				TDCCLCVAGHEFQAHKAILAA
						RSPVFSAMFEHEMEESKKNPSL
						KSMDV\EPEVFKE\MMCFIYTG
						KAPNLTKMA\NDLLAAADKYA
						L\ERLKVMCEDALCSNPV/SVEN
						AAEILIL\ADLH\SADQLKTQAV
				}		DFINYHA\GDVLETSGWKSMVL
						SHP/HNLVAEAYRSLAS\KQCPF
						LGTPTQNALKQSLKILLCC
23705	54073	A	23840	1	575	FRHRKIIY*NLLKQEDSINVTEM
						TSIRASVDILIPDPPPKDDEMDT
						DKQEKKEVPKCGFLPGNEKVL
				į		SLLAMVKPEVWTLKEKCIVVIT
						WIQHLIPKIEDGNDFGVAIQEK
						VLERVNAVKTKVEAFQTTIS/K
						MDYRALVHERDEAAYGELRA
						MVLDLRAFYAELYHIISSNLEKI
						VNPKGEEKPLMYPTRD
23706	54074	Α	23841	1	1092	
23707	54075	Α	23842	130	397	EGVHSQVNLFPIKGNQTHVPAS
		l				HGTDSALPPTLQMDYRALVHE
						PD*AAYGELRAMVLDLRAFYA
						ELYHIISSNLEKIVNPKGEEKPF
23708	54076	A	23843	20	334	VRLPGSTHAGGDRSSMAKPCG
						VRLSGEARIQVEVLRQNLFQEA
						\EEFLYRFLPQKIIYLNQLLQEDS
						LNVADLTSLRAPLDMPIPDPPP
						KDDEMETDEQEKKEGK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
23709	54077	A	23844	1	898	MKQKVAAEAGRGQILQAKVLP SGTLSRLAEALFGVQTYDQDA YGFLTPSAQSMKVFCCLWKLV CKQLEGDVAQGLAIDGDVKKH SGLEWQVETIQNLVDSYMAIV NKNMQDLMPKTIIHLMINNTKE FIFSELLANLYSRGEQNTLMGE SAEQAQGRNEMRRMHHVLRE APSIIGDIKTTIVSTPTGSPKSSPT TQCRLPAVPPARPGLRSPAPGPL PAASTLGEAPVVPSTPGASPDPL
						GPPP\QVPSRPSRGPQPQDRMPC GLAGDLTVGFLEPTLQAIIRWS
23710	54078	A	23845	99	429	LLLRPPANSPESQS FSDAAAVARQPSFGKALGAPQP
						ADTQPTPSMPPRCPGGSETKPK KAAGKDKSSDK\KNKTKKQMK AEKGGKEKQAEVADQETKDLS AENGDTENKESPASDEATEKDT KSD
23711	54079	A	23846	1	390	TWMFIAAVFRIASTWKQPRYFS VGKQDQKRTSRKLQQTCSRGA *WLEGKLTNRKE*HQYQRKGH PRKNTIQRSPTLKTKANIILNGQ KLEAFPLKTGTRQRCPLSPLLFN MVLEVLARSIRQEKEIKVFK
23712	54080	A	23847	522	733	ASAADTQANRVWSGPP/RKLQ QTCS*GS*LLQGKLTNRKDIHT KTPSVHHHHQRPKTLNQQRSK ETKKAIT
23713	54081	Α	23848	46	330	
23714	54082	Α	23849	1	873	MAEKKDPTKEPEKKHLLAWM NKLLYELDPKYKNDSNTDGSA VVHKIIIACDKPVSPPPSAMIGSF PRPSSEADAGAMLCVQPARTKI LGILGLKATSLQYLRLHTGKPL LPKIDESTKMGRNQPKKAENSK NQNASSPPKDHNSSPVREQNR MQNEFDELTEVGFRRDRSSKR NSVVIKALPRHLINQGRSQEWR LGIGARDSPKTSGRRGKSPAGP NRTTVPEDEQA/GSRPRPIPTRQ ADPRTPPPTRSEKE/AQLSLATD QMCPSLAPLIK*PQES**WKQRI SI*MEHL
23715	54083	A	23850	1075	1350	AIITANVFKFVSIDRGLVCNMEL EGSNCFIDVCSWTSEMKSKGTS RR*KIT*GNSSGSLPRDTQRSGH RNSSLCCERSKYGGSRVDIRRK VF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23716	54084	A	23851	1	956	MEVLARVFRQEKEMKGIHIGK
					;	EEHKLFLFADNMTLCLEKTKDS
						VKKLVEQPSSFGSVDWLCLSSC
						SGSTHTPRPADVSPGSLPGPGQ
		ŀ				TFGTREPPQAVSTKEASSSNLH
						APERTVAGLTFTTEQVRALEGV
						FRHHQYLGPLERNWLAREMQL
						SEVQIKTWFQNRRMKHKRQM
						QDSQLNGPLSGSLHGPPAFHSP
						SSGLANGLQLLCPWAPLPGS/L
						WLPLGSFWGLRQMDQEALASS
						WASCCGQPLAYHPPRPGSGAH
1						KLGPALSTGPRGLCALPETGDA
1						FEETPPTTCLPHKRSVQIASDAY
						PEDSAVLFTSWRHLSP
23717	54085	Α	23852	1	846	MEAFDANYTFQSGPSAAGLLV
		1				FAGGPLQALCAWVSAVEAAEQ
ĺ						WILQPSSFGSVDWLCLSSCSGST
						HTPRPADVSPGSLPGPGQTFGT
						REPPQAVSTKEASSSNLHAPER
						TVAGLTFTTEQVRALEGVFRHH
						QYLGPLERNWLAREMQLSEVQ
		1				IKTWFQNRRMKHKRQMQDSQ
						LNGPLMGSFPGAPWLSTHHFLA
						LAKGLQ\LLCPWA\PLPGSPGCP
		1				LGSFWGTP\QMDQEALASSWA
						GCCG\QPLAYHPPRPGSGIDLPK
						GTQLSLRLFQDISSTRGDHRSG
23718	54086	A	23853	3	239	
23719	54087	A	23854	10	1154	RRAGFGGVRASGAMGTPPGLQ
		1				TDCEALLSRFQETDSVRFEDFT
		1				ELWRNMKFGTIFCGRMRNLEK
1						NMFTKEALALAWRYFLPPYTF
						QIRVGALYLLYGLYNTQLCQPK
	1					QKIRVALKDWDEVLKFQQDLV
						NAQHFDAAYIFRKLRLDRAFHF
						TAMPKLLSYRMKKKIHRAEVT
						EEFKDPSDRVMKLITSDVLEEM
						LNVHDHYQNMKHVISVDKSKP
						DKALSLIKDDFFDNIKNIVLEHQ
		1				QWHKDRKNPSLKSKTNDGEEK
						MEGNSQETERCERAESLAKIKS
						KA\FSVVIQASKSRRHRQVKLD
	1					SSPLDSASGQGQVKATRKKEK
						KERLKPAGRKMSLRNKGNVQN
						IHKEDKPLSLSMPVITEEEENES
L	<u> </u>		<u> </u>	<u> </u>	L	LSGTEFTASKKRRQH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23720	54088	A	23855	16	447	VNCDKANCSTTCFNGGTCFYP GKCICPPGLEGEQCEISKCPQPC RNGGKCIGKSKSVCEPGCGAH GTCHEPNKCQCQEGWHGRHCN KRYEASLIHALRP\AGAQLRQH TPSLKKAEERGDPPESN\YIWVE LPTSGNVLSYTKFI
23721	54089	В	23856	103	1989	
23722	54090	A	23857	3	1295	SGRGVSRHMADYLISGGTGYV PEDGLTAQQLFTGAAARDRLT YNNFLILPGFIDFIADKVDLTSA LTWKITLKMPLISSPMDAVTEA DMAIAMALMGGIGFIHHHCTPE FQANEVRKVKKFERGFIMDPV VLSPSHTVG\DVLEGKMRHGFS G\IPITETGTMGSKLDSHKQLLC GAAVGTREDDKYRLDLLTQVG VDVIVLDLSQGNSLYQMAMVY YIKQKYPHLQVIGGNMVTAAQ AKNLIDAGVEGLHVGMGCSSIC ITQEVMAYSWPQGTAVYKVAE YARHFDHGEHGKALALGASTV MMGSLLAATTEVPSKYFFSDG VWLKKYRGMGSLDALEKSSGS QKQYFMEGDKLKIAQGVSSSIQ DKGSIQKFVSYLIAGIQHGCQD TGVCNLSVLLSMMYSGERKFE KWTMLAQIKGGVQGLHSYKK
23723	54091	A	23858	770	1008	LAPSPCVRSKGRKESPDSR/RPL WLRPIAPALGPSGRRGLAPDSR LNPHRSLLGTGNYDVNVIMAA LQGLGLAAVWWDRRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23724	54092	Α	23859	1	1489	MTRWSSYLLGWTTFLLYSYESS
						GGMHEECVFPFTYKGSVYFTCT
						HIHSLSPWCATRAVYNGQWKY
						CQSEDYPRCIFPFIYRGKAYNSC
						ISQGSFLGSLWCSVTSVFDEKQ
	:					QWKFCETNEYGGNSLRKPCIFP
						SIYRNNVVSDCMEDESNKLWC
						PTTENMDKDGKWSFCADTSNL
i						GMGVGPTPRHHRRASTRHSRPP
						PADPDGGGERPRTADDRQDDA
į						TEDDGTRCRGRQTRTQATNAN
						ANHKKSNRTKPPPRHKPNKAK
						RHTKKETPQNT/CGTQDKNPDS
						GRPKLFRRACITSTGTRGELWT
						RSREHPAK\QGRRSGKQRLRPW
				İ		DRMLSPDWHRAQRIAQVWKPE
į						QEDGLSIGFGDQPGPHSDTSSR
						QQVRRSKATEMHVGGRVDFD
						DFVELMTPKLLAETAGMIGVQ
						EMRDAFKEFDTNGDGEITLVEL
						QQAMQRLLGERLTPREISEVVR
						EADVNGDGTVDFEGDIDGAWG
						RHGTQGPHWPWGLGDVRGEPL
23725	54093	A	23860	283	490	GDIWMGTQPKPYHS\PPGPSPN\
						PCPPHFKNQSMPSPKSPPKVFNS
						FPALTQKVHTSKVSSETRQVPS
						HL
23726	54094	С	23861	82	222	
23727	54095	Α	23862	27	454	VDGRIAGEITRRGSRARPRPGPQ
ĺ						CPPGRPGTAMIKAILIFNNHGKP
ĺ						RLSKFYQPYSEDTQQQIIRETFH
						LVSKRDENVCNFLEGG/LVFVE
						TLDKCFENVCELDLIFHVDKVH
						NILAEMVMGGMVLETNMNEIV
						TQIDAQNKL
23728	54096	Α	23863	1	403	RGSRARPRPGPQCPPGRPGTAM
ĺ						IKAILIFNNHGKPRLSKFYQPYS
ĺ		ŀ				EDTQQQIFRETF/HLGSKKEEKG
ĺ						WNFLKGGLLIGGSDNKLDYRH
						YATVYFVF\GGDFSKRELGFLN
i	1	ı	1			LIQEFGEPLDKCFEKVWELDLIF
	54097		23864	215	599	EIQEI GEI EDREI ERV WEEDEN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23730	54098	A	23865	2	683	GRIAGEITRRGSRARPRPGPQCP PGRPGTAMIKAILIFNNHGKPRL SKFYQPYSEDTQQQIIRETFHLV SKRDENVCNFLEGGLLIGGSDN K\LIYRHYAT\LYFVFCVDSSE\S ELGILDLIQVFV\ETLDKCFGNV\ CEL\DLIFHV\DKV\HNIL\AEMV MGGMVLETNMNEIVTPIDAQN KLEKSEAGLAGAPA\RA\VSAV KNMNLP\EIPRNINIG\DISIK\VP
23731	54099	В	23866	94	726	
23732	54100	В	23867	94	1146	
23733	54101	В	23868	1	360	
23734	54102	В	23869	1	1371	
23735	54103	В	23870	1	639	
23736	54104	В	23871	280	2391	
23737	54105	A	23872	3	244	
23738	54106	В	23873	103	285	
23739	54107	A	23874	70	378	MPLRILLMHGEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE
23740	54108	A	23875	5	156	RKKGGG/EREEEGEGEEEEEE EEEEEEEEEEEEEEEEVVL LTPGTGAEDFKKH

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SEQ ID	~		SEQ ID NO:	i		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23741	54109	Α	23876	1	1808	MEVLSIDNLAGHKVFYQKESHS
						WKQQKQDLMSGKRLSAANPLP
						ALLLPVLWTTGQAPSPGPGGDP
						ILQLWDDAQQMAGSPRTKKKG
						CTRNSQAARLSFQSRGGQGSLT
						EKMIFEQRFGGGKGESRKDAIA
						IIVVDIVTLQPCRTVEWVPTRPF
						RSLQGHFQGGLRRQQMMVME
						GVWVLSCAMDLTSPRDCLPFC
ļ						VSLPVLLLSAPLCSCSTAGGSPV
						TSPHHFPVEGGLGRVRRCYINE
						PVVTTLGWESVCWVSFWLLLR
					İ	IRYSTSGQAACHAESQELNPRG
						FGCLGHRNVTWVKCLLPALQP
		1	}			EGSVPFLELLEPFIFASASKTFV
						AANLVTWVACFLLVQAAFQLS
						EYRQGVGIPVADRTGDAPTLGL
		1				SYEADVAEGHGFFLDQESVIQII
						PEPLPQFYLCLLDAELTGVQGE
						PALPSMQIFSSPEAATGVHNDR
						GAEHFPVSRQPSAELGLRPMQA
						SQERKINFPGPSPGFPPKRQNKT
						LSRREKKEEEKEEKEEEEEE/E
						GEEEKEKEEEEEEEE
						GGEEEEEEEEEEEEEE
						KWLYQYDPEDKAQSKQWLSR
						GGSGPVKAKADWWRAKEIHSL
		<u> </u>				QPRQVIPGIPSM
23742	54110	Α	23877	459	636	IPGGSNGLEAEGPPPPPLPVAAV
						LA/AGSGGLAVASCAFTICALR
22-12			22272	100		WPGVPCSRVPIWTPS
23743	54111	C	23878	183	254	L COCOL DWED IDLOCCOL LIER
23744	54112	Α	23819	103	335	LGSGDLPWEINPLSSCSLLHEK DPPTTSGPQTDRPKEHLINFKSG
						ACYTCRKSGH*AKECLQPGIPIK
						PHLICVGPH
23745	54113	A	23880	159	306	LGSGNLP*EINPLSSCSLFREEDP
23743	34113	Γ.	23660	139	1300	PTTSGPQTNQPKEHLTNFKSAA
						ED
23746	54114	С	23881	80	106	
23747	54115	A	23882	317	427	
23748	54116	A	23883	353	511	
23749	54117	В	23884	1	320	
23750	54118	A	23885	1	624	
23751	54119	A	23886	224	355	
23752	54120	A	23887	872	1003	
23753	54121	A	23888	1	603	
23754	54122	A	23889	2	214	
23755	54123	A	23890	459	601	DVDRHVRGSNFHHNEIRSLAAT
		-			-	PRAPGTLAQ/GLTDSFPDLLGLA
l	Ī	I	l	I	1	AED

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23756	54124	A	23891	2	267	QWDGWGARKGMEWEDDLPLE FGRPTASLLSDRPQPNSSRGSDV PFL\PPSLPCHSSACLAATPRAPG TLAQGSLTDSFPDLLGLAAED
23757	54125	Α	23892	72	299	
23758	54126	В	23893	1	1470	
23759	54127	A	23894	134	1276	
23760	54128	Α	23895		847	PPPPRVLRTGGGASLTAQAPGT PRRPQEGEPVTPGRSLGLRSSCQ ASVSLLLPVIVACASLEPGRGR AFRDEGAWNTRVSAAVISSQA QLLSSILVNSRRLSFSAHHTAFG TVALGEYAFRMF\RLSARLFG\E VT\RPTNS\RSLMKVVKLFSE\IW PLAKKKETYDW\YPNHHTY\AL ELMAGRLRFLGTLTEDEAFRDF MGMEAKTDLKEAFVGKGGNP KEREKGKRAAKKGNSCLVPSR RGDFLSSVAERRKLHLLSFPHIG GDVIFLNGSLFGRNTSHLLGEI
23761	54129	Α	23898	3	1224	MRRNKPRQRGVREARPAGGAG PGWRGARCSGAGEGGGGERRG SPPAALALAPASGPRRNFPPDA RCLIYQQIHQGAFLLAGSASLS AVTGVPEGEARFTEDYCPEEK MFGFHKPKMYRSIEGC\CISGA KSSSSRFTDSKRYEK\DFQ\SCFG LHETRSGDI\CNACVLLVKRWK KLPAGSKK\NWNHVVDARAGP S\LKTTFEPK\KVKTLSG\NRIK\S TQISKLQKEFK\RSNSDAHSTTS SASPAQSPCYSNQSDDGSDTEM ASGSNRTPVFSFLDLTYWKRQK ICCGIIYKGRFGEVLIDTHLFKP CCSNKKAAAEKPEEQGPEPLPIS TQEWTECPDLSLVVSWRPPLSS LPAGCSLHGLAADWTTDMITV TTSKCREPSCQQEGSFQRRKAF PKRLPKMAEVQVLVL
23762	54130	Α	23900	69	355	
23763	54131	Α	23901	316	830	VSLSQTLHPPMTLQLSRYSRGW GLSSDRPRGAKAKVPSCFGTAP R/GPEAAPRLLAHHLVPAGRGS/ ARAPAPGWRLGSP\GAGRKGR DFRLPASGLAVSGEWDPIPWGY VGEVGGCSCQWFSPGVQQKRR RRKVWLGQPQDLPGPSALADK GPALAKADGQRVNVASVGLQP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23764	54132	A	23902	1	1044	MSYKVTQEISKELRWISAVLVT
						WDLRGETGTRVCIAMGVSRMQ
						YSIMAKIVGSGNNLELNPAADK
						LDNLGAQCGLASVKVSARAAP
						PFAPATLVALLEARAAELPLPPP
						PGVSLRCVEAPDVRGRGRGGG
						EGSGVTRLLLPPPPPSGASFLQR
						GRPVTPALFQAGWGRAADRPG
						PADLAGGAGGRSGGRRMLPRS
						PGAALRDRSALEDGIVEGYGPL
						GEPEEGSRSSFLESKVAFNYCN
			ļ			KVAPLDFGNEAVEQCHTMSDR
						KAVIKNADMSEDMQQDAVDC
						A\TQAMEKYNIEKDIAAYIKKEF
						DKKYNPTWHCIVGRNF\GSY\V
						PHTEDKHFIYFYLGQVAIL\LFK
23765	54133	Α	23903	33	427	
23766	54134	Α	23904	72	777	EWKQAGEGGGGQRRGSPPAVL
						ALAPASCPRREMSYLPTNPPGSI
						LICWLGVSLSSHRSAGGGSAIY
		İ				G\SYYPEEKMFVFHKPKMYRSI
						EGYCICRAKSSSSRFTDSKRYEK
						DFQSCFGLHETRSGDICNACVL
						LVKRWKKLPAGSKKKTGIMCN
						QSDDGSDTEMASGSNRTPVFSF
						LDLTYWKRQKICCGINYKGRF
						GEVLIDTHPFKPCCSNKKAAAE
						KPEEQGPEPLPISTQEW
23767	54135	Α	23905	234	989	PKTSDQAYYCPEEKMFGFHKP
						KMY\RSIEGC\CISGAKSSSRFT
			i			DSKRYEK\DFQ\SCFGLHETR\SG
						DI\CNACVLLVKRWKKLPAGSK
						K\NWNHVVDARAGPSLKTTLK
						PKKVKTL\SGNRIK\STQISKLQK
						/ENFKR\HNS\DAHSTHPQVPSP\
						AQSPLFTVNQFRWTGSDTGVG
						FPGSNRNHPVFSFLDL\TYWKR
						QKICCGI\IYKGRFGEVLIDTHLF
						KPCCSNKKA\AAEKPEEQGPEP
						LPISTQEWVTEVFM
23768	54136	A	23906	3	1059	

NO:	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23769 54137 A 23907 1 1697	NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
23769 54137 A 23907 1 1697		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
ESRGGAPDTPRRLHSFTQTRSLS PQAHAGPHTRLHPRQGHAHM GGRVTDTRRHIPNADVPRYK LTCTGGVTTDTRHIPNADVPRYK LTCTGGVTTDTRHIPNADVPRYK LTCTGGVTTTVALYDYESRTET DLSFKKGERLQIVNNTKGDW WLAHSLSTQGTGYIPSNYVAPS DSIQAEEWYFGKITRRESERLLL NAENPRGTFLVRESETTKGAYC LSVSDFDNAKGLNVKHYKIRK LDSGGFYITSRTQFNSLQQLVA YYSMSHCPPFSAAGWYGGGCF EEPQKHADGLGHRLTTVCPTSK PQTQGLAKDAWEIPRESLRLEV KLGQGCFGEVMGTWMGTTITT VAIKTLKRGTIMSPFALQEAQV MKKLRHEKLVQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLJDMAAGJASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PCCPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PCCPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PCCPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PGCPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYKSTURE PCCRC PCCPC PCCPESLHDLMCQGRYSTURE PCCRC PCCPC PCCPESLHDLMCQGRYSTURE PCCRC PCCPC					sequence		
PQAHAGPHTRLHPRQGHAHMR GGGRYTDTRRHIPNADVPRYY	23769	54137	A	23907	1	1697	MRPCALCWRSAVTAAPERAVL
PQAHAGPHTRLHPRQGHAHMR GGGRYTDTRRHIPNADVPRYY							ESRGGAPDTPRRLHSFTOTRSLS
GGGRVTDTRRHIPNADVPRYK							1 ` 1
LICTGGVTTFVALYDYESRTET							1 '
DLSFKKGER\LQ\IVNNTKGDW WLAHSLSTGQTGY PSNYAPS DSIQAEWYFGKITRRESERLL NAEMPRGTFL\VRESETTKGAY\C LSVSDFDNAKGLNVKHYKIRK LDSGGFYITSRTGFNSLQQLVA YYSMSHCPPFSAAGWYGGGGP EEPQKHADGLCHRLTTVCPTSK PQTQGLAKDAWEIPRESLREV KLQGGCFGEVWMGTWNGTTR VAIKTLKPGTMSPEAFLQEAQW MKKLRHEKL\VQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQL\DMAAQIASGMAY\V ERMNY\HRDLRAANIL\VGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAAL\VGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAAL\VGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAAL\VGENL VCKVADFGLARLIEDNEYTAR QFAKFPIKWTAPEAAL\VGENL VCKVADFGLARLIEDNEYTAR VGKVADFGLARLIE							
WLAHSLSTGQTGYIPSNYVAPS DSIQAEEWYFGK1TRRESERLLL NAEMPRGTFLVRESETTKGAYO LSVSDFDNAKGLNVKHYKIRK LDSGGFYITSRTQFNSLQQLVA YYSMSHCPPFSAAGWYGGGCP EEPQKHADGLCHRLTTVCPTSK PQTQGLAKDAWEIPRESLILEV KLQGCFGEVWMGTWNGTTR VAIKTLKPGTMSPEAFLQEAQV MKKLRHEKLVQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMMYVHRDLRAANILVGENL VCKVADFGLARLJEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHOLMQCWRKPEER PTFEYLQALEDDEYTSTEPQVQP PECPESLHOLMQCWRKPEER PTFEYLQALEDVFTSTEPQVQP PECPESLHOLMQCWRKPEER TFCEACKSTVGVDFKIKTVELR GKKIRLQIWDTAGQERSSITSA YYRSAKGMILVYDITKKETFDD LPRWMKMIDKYAISEDADLLL VGNKLDCETDREITRQQEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPRRPHVR 23771 54149 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23914 3 481 23777 54144 A 23914 3 481 23777 54145 A 23915 I 711	1						
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NAENPRGTFLVRESETTKGAYC							,
LSVSDFDNAKGLNVKHYKIRK LDSGGFYITSRTQFNSLQQLVA	ŀ						l -
LDSGGFYITSRTQFNSLQQLVA							
YYSMSHCPPFSAAGWYGGGCP EEPQKHADGLCHRLTTVCPTSK PQTQGLAKDAWEIPRESLRLEV KLGQGCFGEVWMGTWNGTTR VAIKTLKPGTMSPEAFLQEAQV MKKLRHEKLVQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPCA PCMVNREVLDQVERGYRMPC	İ						l l
EEPQKHADGLCHRLTTVCPTSK PQTQGLAKDAWEIPRESLRLEV KLQQGFGEVWMGTWNGTTR VAIKTLKPGTMSPEAFLQEAQV MKKLRHEKLVQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADPGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP ADVINGSRGVGKTSLMERFTDD LPKWMKMIDKYALSEADLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23777 54145 A 23915 I 711	}						
PQTQGLAKDAWEIPRESLRLEV KLGQGCFGEVWMGTWNGTTR VAIKTLKPGTMSPEAFLQEAQV MKKLRHEKLVQLYAVVSEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLIEDNEYTAR QGAKFJIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQVQP LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGVDFKIKTVELR GKKIRLQIWDTAQGERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYAISEDADLLL VGNKLDCETDREITRQQEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPPPHVR 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23913 3 143 23776 54143 A 23913 3 1443 23777 54145 A 23915 I 711	1						1
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VAIKTLKPGTMSPEAFLQEAQV MKKLRHEKLVQLYAVVSEEPI YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQC WRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PGCPESLHDLMCQC WRKEPEER PGCPESLHDLMCQC WRKEPEER PGCPESLHDLMCQC WRKEPEER PGCPESLHDLMCQC WRKEPEER PGCP							l · ·
MKKLRHEKLVQLYAVVSEEPI							l
YIVTEYMSKGSLLDFLKGETGK YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP							1
YLRLPQLVDMAAQIASGMAYV ERMNYVHRDLRAANILVGENL VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PGMVNREVLDQCWRKAPCH PTFEYLQAFLEDYFTSTEPQYQP PGMVNREVLDQCWRKAPCH PGMVNREVLDQCWRCHAPCH PGMVNREVLDQCWRAPCH PGMVNREVLDQCWRCH PGMVNREVLDQCWRCH PGMVNREVLDQCWRCH PGMVNREV							1
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VCKVADFGLARLIEDNEYTAR QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP							I
QGAKFPIKWTAPEAALYGRFTI KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPEER PTFEYLQAFLEDYFTSTEPQYQP 23770 54138 A 23908 I 736 MDPGAALQRRAGGGGLGAG SPALSGGQGRRKQPPRPADFK LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGVDFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 I 711							
KSDVWSFGILLTELTTKGRVPY PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQC WRKEPEER PTFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQP PFEYLQAFLEDYFTSTEPQVQPRADFK LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGVDFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYANSEDADLLL VGKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI PLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PRIVATE PKLVDDILKKMPLDILRNELS PKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKKMPLDILRNELS PKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILRNELS PKLVDDILKMPLDILKMPLDILRNELS PKLVDDILKMPLDILKMPLDILRNELS PKLVDDILKMPLDILKMPLDILKMPLDILKMPLDILKMPLDILKMPLDILK							
PGMVNREVLDQVERGYRMPCP PECPESLHDLMCQCWRKEPER PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQP PTFEYLQAFLEDYFTSTEPQYQFRAGGGGGLGAG SPALSGGQGRRRKQPPRPADFK LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGV\DFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR PTFEYLQAFLAGA PTFEYLQAF							1 -
PECPESLHDLMCQCWRKEPER PTFEYLQAFLEDYFTSTEPQYQP 23770							
PTFEYLQAFLEDYFTSTEPQYQP 23770 54138 A 23908 I							·
23770 54138 A 23908 I							l
SPALSGGQGRRKQPPRPADFK LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGV\DFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR							<u> </u>
LQVIIIGSRGVGKTSLMERFTDD TFCEACKSTVGV\DFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 I 711	23770	54138	Α	23908	1	736	· · · · · · · · · · · · · · · · · · ·
TFCEACKSTVGV\DFKIKTVELR GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711							i
GKKIRLQIWDTAGQERFSSITSA YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711			1				LQVIIIGSRGVGKTSLMERFTDD
YYRSAKGMILVYDITKKETFDD LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR							TFCEACKSTVGV\DFKIKTVELR
LPKWMKMIDKYA\SEDADLLL VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711			l				GKKIRLQIWDTAGQERFSSITSA
VGNKLDCETDREITRQQGEKFA QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR							YYRSAKGMILVYDITKKETFDD
QQITGMRFCEASAKDNFNVDEI FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPPRPHVR 23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 I 711			}				LPKWMKMIDKYA\SEDADLLL
FLKLVDDILKKMPLDILRNELS NSILSLQPEPEIPPELPPRPHVR 23771 54139 B 23909 84 441			I				VGNKLDCETDREITRQQGEKFA
NSILSLQPEPEIPPELPPRPHVR							QQITGMRFCEASAKDNFNVDEI
23771 54139 B 23909 84 441 23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711			l				FLKLVDDILKKMPLDILRNELS
23772 54140 A 23910 6 254 23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711							NSILSLQPEPEIPPELPPPRPHVR
23773 54141 A 23911 1 230 23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711		<u> </u>	В			<u> </u>	
23774 54142 A 23912 3 187 23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711		54140	Α		6		
23775 54143 A 23913 3 143 23776 54144 A 23914 3 481 23777 54145 A 23915 1 711			Α				
23776 54144 A 23914 3 481 23777 54145 A 23915 1 711			Α		3	187	
23777 54145 A 23915 1 711		54143	A	23913	3	143	
	23776	54144	A	23914	3	481	
23778 54146 A 23916 55 381	23777	54145	Α			711	
	23778	54146	Α	23916	55	381	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
02770	64147	<u> </u>	100017	1040	1076	CL CA POTRITON ACAMELON ATTACK
23779	54147	Α	23917	240	875	GLSAESTPTSIMSMTLGYWDIR
			1			GLAHAIRLLLEYTDSSYEEKKY
						TMGDAPDYDRSQWL\NEKFKL
						GL\DFPNLPY\LIDGAHKMHPEA
						TPFLCYI\ARKHNLLLGETGRRE
						KISSWDIL\ENQAMDVSNQLAR
						VCYSPDFEKLKPEYLEELPTMM
						QHFSQFLGKRPWFVGDKITFVD
						FLAYDVLDLHRIFEPNCLDAFP
22500	711.10	<u> </u>	22212			NLKDFISRFEVSCGIM
23780	54148	Α	23918	64	774	CLSAESAPTSTMPMTLGYWDIR
						GLAHAIRLLLEYTD\SSYEEKKY
1						TMGDAP\DYDRSQWLNEKFKL
		1				GL\DFPNLPYLIDGAHKITQSNA
						ILRYI\ARK\HNLCGGDRRREKIS
						CWDILENQVM\DNHMELVRLC
						YDPDFEKLKPKYLEELPEKLKL
						YSEFLGKRPCFLG\DKITFVDFL
						VY\DVLETK\QVFEPK\CLDAF\Q
			İ			NLKDFISRFEGL\EKISAYMKSS\
						RFL\QELVFSKMAVWGNK
23781	54149	Α	23919	163	317	LSPNNYYRKLPQKVKLE*RYHR
-						LALFSFPTKKLTPEKVSLCCRFL
		<u> </u>				EMGYIF
23782	54150	Α	23920	62	255	VRLTAQKSLGPLTSTHSLWGSA
						\RARHREPVGSVCGCDPCEQLL
						LLVSQLRAPGVDSAAAGRPV
23783	54151	A	23921	173	467	KSRFCFFGVVLKLPKKLHIR*WI
						GATL*NQP*TIFQTDGDCSHQSL
						GPLPACDLWDQLHLRSKQGGS
						VCGCDPCEQLLLLVSQLRAPGV
	ļ	_		<u> </u>		DSAAAGRPV
23784	54152	A	23922	35	548	PRSCPSLSPGLSCLLQ/RLVNYQI
	1	1				SIKCSNQFKLEVCLLNAENKVV
						DNQAGTQGQ\LPAVGQARWGS
						GCRGAHTW\PLAVIGAADCTG
						QRGWEGV*AFPGLTEPVSSLFC
						GQDGD/CSPHSM*VLSHVVLWL
						LNGLC\SLVLGSVCGCDPCEQL
						LLLVSQLRAPGVDSAAAGHPV
23785	54153	Α	23923	1	599	MSPAGLVNYQISVKCSNQFKLE
		1				VCLLNAENKVVDNQAGTQGQL
						KGRPDGAQAV/VGAHTWRFGC
						RRCG*LHRSHWGL*LLHTPCGR
						PHCARHREPVAHSLLCSRAISGS
						EPHFWQCSVQ*PQLCVPGRVAT
[1				PLGELDSGDGDCSHQSLGPLPA
1			1			CDLWDQLHLRSRQGGSVCGCD
		Ì				PCEQLLLLVSQLRAPGVDSAAA
						GRPV
23786	54154	Α	23924	290	528	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23787	54155	A	23925	274	571	KSRFCFFGVVLKLPKKLHIR*WI GATL*NQP*TIFQTDGDCSHQSL GPLPACDLWDQLHLHSRQGGS VCGCDPCEQLLLL\VSQLQAPAI DSAAAGRPV
23788	54156	В	23926	204	410	
23789	54157	A	23927	29	330	YLGVELVKSAYPFFQIHLSFPSC MWLCGLFLLFYIDFLTLMCGD KPLICKKHQLSASTCA/CTIC*G ALTLLMEIEQGPCDLSCASREV STARSLAELFSGP
23790	54158	A	23928	259	699	ETSHQVMDRSNPVKPALDYFL NRLVNYQISVKCSNQFKLEVCL LNAENKVVDNQAGTQGQLKV LGANLWWPYLMHEHPAYLYS WEDGDCSHQSLGPLPACDLWD QLHLRSRQGGSVCGCDPCEQLL LLVSQLRAPG\VDSAAAGHPV
23791	54159	A	23929	245	695	
23792	54160	A	23930	302	745	ETSHQVMDRSNPVKPALDYFS NRLVNYQISVKCSNQFKLEVCL LNAENKVVDNQAGTQGQLKV LGANLWWPYLMHEHPAYLYS WEDGDCFTPKALDPLPACDLW DQL\HLRSRQGGSVCGCDPCEQ LLLLVSQLRAPGVDSAAAGRPV
23793	54161	٨	23931	2	293	VFCHTDLRKNLTVDEGTMKVE VLPALTDNYMYLVIDDETKEA AIVDPVQPQKVVDAARKHG\V KLTTVLTTHHHWDHAGGNEKL VKLESADLGIRLF
23794	54162	A	23932	1	441	
23795	54163	A	23933	1	630	
23796	54164	Α	23934	3	265	
23797	54165	A	23935	3	1062	SAVQPGPVMVVGRGCSRRSLA ALGAACARRGLGPALLGVFCH TDLRKNLTVDEGTMKVEVLPA LTDNYMYLVIDDETKEAAIVDP VQPQKVVDAARKHGVKLTTVL TTHHHWDHAGGNEKLVKLESG LKVYGGDDRIGALTHKITHLST LQVG\SLNVKCLATP\CHTSGHI CYFVSKLGG\SEPFVVFTGDTLF V\AGCGEVLMKGLRDEMCKAL LEV\LGRLPPDTVGSASLRAGR VYCG\HEYTIKQTSKFAR\HVET GNA\AIREK\LPWA\KEKYSIGEP TVPST\LAE\EFTYNP\FIKVREER RCKQH\AGETDPVTTMRPVRRE KDQ\FK\MPRDLRPPLHLQQDL GIKALLDRLF

SEQ ID NO:	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23798	54166	Α	23936	107	750	SHLWSPVSEPLQACSFC/VIRDK
		İ				YRYFACLMRARSEEHTNEKDM
						MKAIRLLKEAEEEFWYHQHPQ
						PCIFPDSPRGTSYVTHECYKVPE
						WCLDDWHPSEKAMYPDYFAK
						RKQWKKLRRESWKREVKQLQ
						EETPPAGKEECKLHEGMNYVV
				-		LTCIPGAIKQYLACSRYSKTPGP
				1		RDSCWEVQSDNQCWWWLDSG
						AHGLVTSFKGVMMTVEATTM
				1		AAVF
23799	54167	Α	23937	3	718	PASTISCSVTRSRRAVSRLSARP
						GKVSAVMASLASGPYLTHQQK
						VLRLYKRALRHLESWCVQRDK
						YRYFACLMRARFEEHKNEQDM
						AKATQLLKEAEEE\FWYRQHPQ
						P\YIFPDSPWGHPPYERY\DCYK
		ļ				VP\EWCLDDWHPSEK\AMYP\D
						YFAKREQWKKLRRESWEREV\
						KQL\QEET\PPGGSF*LEALPPGP
	ŧ					KRKVILPPLWW\YIVTRP\RERP
		1				MLERERPHLSCLQVKYVTEHG
23800	54168	Α	23938	269	488	TDGVIFIGLSWSPRRCGLGIQRD
						AERIFFALDQASCGFTIHPHIDA
						AETPRGLT\MHLCRRCQRGQSQ
						AEKLP
23801	54169	Α	23939	1	1560	MVHLGAPVMALNLQGKHLCD
						STASSLLPRRRESNSDDTARED
						ARQKRPANTDMRTQELQTRET
						GQGTARETEVATPLSGPSAGDS
						PPAASEMSCGQLVLALLLLFFL
						LVWNTGTKSGSGRVARLAQLA
						HDCDGGLALGLCYAVLDQRSV
						VQATPQAPKPGPVQPLTVQGL
						QPVHVAQESSGNEITARRQSAR
						SRPRGDRTAPSRRVTRRPKHAP
		1				ERAPIQQGNIVTVRAPTARKKW
		1				RPAVGTPTGAHPVCTAKGTGG
		1				RPTPTQAHTRYTERDPIRNSAEL
						TGSRPCTGRERSGVEACKAAV HNLL\AQTGSSGLIINYPLDD\Q
						1
				1		DAIAVEAACTNVPALFLDVSDQ
						TPINSIIFSHEDDVWRTPLVAPT
						NLNTRSSQPARHHRAGAVRRH
						HMTVARSATAHARLRRHAKPQ GRERIKPSGAREGSEPLGGESGI
						RTPRGRERDPNPSGAKAGSEPG
						RIPETLPRCKGETEAKCWARGN
						FHTAQGSWPRGGRSRWEPLEA
22000	64170	 	22040	100	220	PALSQQAHGGELPLPAAARSHR
23802	54170	В	23940	98	229	

SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23803	54171	Α	23941	11	256	
23804	54172	Α	23942	289	433	VLWSCAAEDFDWLVYRPIGSA
						LSCRLIGPWAIGMWETGFP*VL
						WSCAAEDFDWLVYRPIGSALSC
						RLIGPWAIGMWETGFPLLSSVS
		1				LS
23805	54173	Α	23943	450	1260	LCACVVGGEKRSGPRIGCAGA
						GKDPFWRAVGVAQKAAAVVA
				j		CGAASPYRNRADVSVSDHSQS
						RGPGGLSLSGGTVCDMGKKTK
		ļ				RTADSSSSEDEEEYVVEKVLDR
						RVVKGQVEYLLKWKGFSEEHN
		ł				TWEPEKNL\DCPELISEFMKKY
		Ì				KKMKEGGN\NKVREKSESSKR
						KSNFSDSAEDIKCKKKREQSND
						IARGFERGLEPEKIIGATDSCGD
		ŀ				LMFLMKWKDTDEADLVLAKE
		ŀ		i		ANVKCPQIVIAFYEERLTWHAF\
						PEDAENKEKETAKS
23806	54174	Α	23944	3	267	AMVGGGGVGGGLLENANPLIY
						QRSGERPVTAGEEDEQVPDSID
]		AREIFDLIRS*AWPLTPQPAYW
				ļ		YPGPSSCQGCYRCFLESLTMRN
23807	54175	Α	23945	1	942	
23808	54176	Α	23946	3	470	AMVGGGGVGGGLLENANPLIY
						QRSGERPVTAGEEDEQVPDSID
						AREIFDIRRCWARAGSGGLRW
						GEQ*YRAAGGAASQQGVPGRG
						F*VGKGAESLFLHFPQ\LIRFLN
						DP\EHSLTLEELNVVEQVRVQV
						SDPESTVAVAFTPTISHWSMGT
23809	54177	Α	23947	2	604	ARSHRISGGGSAMVGGGGVGG
						GLLENANPLIYQRFGE\RPVTAG
		İ				EEDEQVPDSIDAREVF\DLIRSH
						QMTPEHP\LTLEELNVV\EQVR\
						VQVSDP\ESTVAVGFPKPTHSGT
						CRHGPPLIGLS\IKVK\LLRSLPS
			,			AFSRWDVAHLLPGDPLPQEHA
						VEQATLQIKEAGWATALGRTH
						HLLEVVN\QCLSSPAPGTWAFV
23810	54178	Α	23948	1	1002	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	, .
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ĺ				sequence		
23811	54179	Α	23949	89	2587	STSRDRRLSKRSRTTMAAQPRG
ĺ						GDYRGSFLLSILLGTPWEAWAG
		1				RILYSVSEETDKGSFVGDIAKDL
}						GLEPRELAERGVRIISRGRTQLF
f						ALNORSGSLVTAGRIDREEICA
						QSARCLVNFNILMEDKMNLYPI
İ						DVEIIDINDNVPRFLTEEINVKI
						MENTAPGVRFPLSEAGDPDVG
		l				TNSLQSYQLSPNRHFSLAVQSG
		ł				DDETKYPELVLERVLDREEERV
		ŀ		1		HHLVLTASDGGDPPRSSTAHIQ
		l				VTVVDVNDHTPVFSLPQYQVT
						VPENVPVGTRLLTVHAIDLDEG
						VNGEVTYSFRKITPKLPKMFHL
						NSLTGEISTLEGLDYEETAFYE
						MEVQAQDGPGSLTKAKVLITV
						LDVNDNAPEVTMTSLSSSIPED
						TPLGTVIALFYLQDRDSGKNGE
						VTCTIPENLPFKLEKSIDNYYRL
						K\TTKNLDRETLSLYNITLKATD
						GGTPPLSRETHIFMQVADTNDN
						PPTFPHSSYSVYIAENNPRGASI
						FLVTAQDHDSEDNAQITYSLAE
						DTIQGAPVSSYVSINSDTGVLY
						ALQSFDYEQLRELQLRVTAHDS
						GDPPLSSNMSLSLFVLDQNDNP
						PEILYPALPTDGSTGMELAPRSA
						EPGYLVTKVVAVDKDSGQNA
						WLSYLLLKASEPGLFAVGLYTG
		ĺ				EVRTARALLDRDALKQSLVVA
1]				VQDHGQPPLSATVTLTVAVAD
						SIPEVLADLGSLEPSDGPYNYDL
23812	54180	В	23950	165	335	
23813	54181	Α	23951	217	859	GSAMAQQRALPQSKETLLQSY
ļ	,					NKRLKDDIKSIMDNFTEIIKTAK
ļ					1	IEDETQVSRAT\QGEQDNYEMH
						VRAANIVRAGESLMKLVSDLK
						QFLIL\NDFPSVNEAIDQRNQQL
						RTLQEECDRKLITLRDEISIDLY
						ELEEEYYSSSPGPSLVSQQLSEQ
						QLLSSAWTAWPKVTLSSLAPGP
						QYSVNGPAAIIAHPGTCASAGIS
						SSLLPEPVLLVPVF
23814	54182	Α	23952	1	1263	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	į			sequence		
23815	54183	A	23953	70	2530	PGVSRLVSKLRRARGMGSGAG
23013	54103		23733	, 0	2550	ELGRAERLPELFLFLLTLFCPAL
						CEQIRYRIPEEMPKGSVVGNLA
	İ				,	TDLGFSVQELPTRKLRVSSEKP
						YFTVSAESAELLVSSRLYREEIC
1	1					GKKPACALEFEAVAENPLNFY
						HVNVEIEDINDHTPKFTQNSFEL
						QISESAQPGTRFILGSAHDADIG
						SNTLQNYQLSPSDHFSLINKEKS
						DGSKYPEMVLKTPLDREKQKS
						YHLTLTALDFGAPPLSSTAQIH
						VLVTDANDNAPVFSQDVYRVS
						LSENVYPGTTVLQVTATDQDE
						GVNAEITFSFSEASQITQFDLNS
						NTGEITVLNTLDFEEVKEYSIVL
						EARDGGGMIAQCTVEVEVIDE
		i				NDNAPEVIFQSLPNLIMEDAEL
						GTHIALLKVRDKDSRHNGEVT
						CKLEGDVPFKILTSSRNTYKLV
		1				TDAVLDREQNPEYNITVTATOR
						GKPPLSSSSSITLHIGDVNDNAP
						VFSQSSYIVHVAENNPPGASISQ
		1				VRASDPDLGPNGQVSYCIMAS
						DLEQRELSSYVSISAESGVVFA
						QRAFDHEQLRAFELTLQARDQ
						GSPALSANVSLRVLVDDRNDN
						APRVLYPALGPDGSALFDMVP
						HAAEPGYLVTKVVAVDADSGH
	1	ŀ				NAWLSYHVLQASEPGLFSLGLR
						TGEVRTARALGDRDAVRQRLL
						VAVRDGGQPPLSATATLHLVF
		1				ADSLQEVLPDITDRPDPSDLQA
23816	54184	С	23954	75	149	
23817	54185	A	23955	442	618	CQRLRSWVSQPFSFSFFPSFPTF*
		` `		~		NILCLFVYLEVEFPSCCPGWSN
						NWAKSSGKALEAN
23818	54186	Α	23956	1	245	
	15 . 1 . 0	<u>'``</u>	1-2220	<u> </u>		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23819	54187	Α	23957	30	11079	APCASFLTPLAAASVGKVWLG
						LRGAAPMASVLSYESLVHAVA
						GAVGSVTAMT\VVFPLDTASIF
İ					ļ	DFSVDF\KRKSKTTHMVVLEIIK
		1		1	•	EEGLLAPYRGWFPVISSLCCSNF
						VYFYTFNSLKALWVKGQHSTT
						GKDLVVGFVAGVVNVLLTTPL
						WVVNTRLKLQGAQFRNEDIVP
						TNYKGIIDAFHQIIRDEGISALW
						NGTFPSLLLVFNPAIQFMFYEGL
						KRQLLTFAFQLSSLDVFIIGAVA
						KAIATTVTYPLQTVPLFPQFGR
						HRLNPENRTLGSLRNILYLLLQI
						FRTPSEREPFYTSVTLSVLSSVA
						YPMMPLSNITSAYRIGVLEKGG
		1				GSGKSDNSDDDDDDDDDGGV
23820	54188	Α	23958	1375	2451	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE\
		ŀ				AEEERVVDNRMHDRFCELRTA
		l				AGHPVEGGRPLVIKGITHPAET
						ALQRFKFLMYQVVASDLDGTL
						LSPDHTLSPYAKETLKLLTARGI
						NFVFATGRHHVDVGQIRDNLEI
						KSYMITSNGARVHDLDGNLIFA
		ŀ				HNLDRDIASDLFGVVNDNPDIIT
						NVYRDDEWFMNRHRPEEMRFF
						KEAVFQYALYEPGLLEPEGETK
·						ALLTPGKLKNFLVEKMRTLGT
						AACPPYHIAFVIGGTSAETNLKT
						VKLASAHYYDELPTEGNEHGQ
[GSAMSSWNRNCWKRPRNSVLA
	}	1				RSLAVNTSRTTFALSVCHVTAH
		<u> </u>			_	PARSAWASPVR
23821	54189	В	23959	1	1386	
23822	54190	Α	23960	1375	1609	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE
						S\EEERVVDNRMHDRFCELRTA
	<u> </u>	<u> </u>				AGHPVEGGRPLVIK
23823	54191	A	23961	433	741	
23824	54192	A	23962	12	224	
23825	54193	Α	23963	3	2053	

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NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23826	54194	Α	23964	1	2408	MGGADIAGPRREHNAKHPTVG
						LGSAAQTTCLPGPKQKGDPTRH
}						QPVDKRNDVGPRQHFWCWRV
		<u> </u>				EIHKEKRMKTGPLNESELEWLD
						DILTKYNTDHAILDVAELDGLL
						TAVLSSPQEIEPEQWLVAVWG
						GADYVPRWASEKEMTRFMNL
						AFQHMADTAERLNEFPEQFEPL
						FGLREVDGSELTIVVPPGGRTA
						TNALSCGSLFGEETSPRGPSADS
						TSQSIGGGSPHDCAKGIALVAA
	ļ ļ					NGGDIRDYEGVDRSAKPQLPMI
						AINTTAGTASEMTRFCIITDEAR
1	ļ					HIKMAIVDKHVTPLLSVNDSSL
						MIGMPKSLTAATGMDALTHAI
						EAYVSIAATPITDACALKAVTM
						IAGISTFFASSRMALMQASAPSA
						SFLPVTFTPIAAAQSRRRAAAT
						GYRDRWLCHGDVLLQICANTQ
						DTVIHALRDIIKHNSCPDDCSVQ
		l				HDAVIDDGCDFIDYLGGSETVV
						ALNTPRADDYHPVMVLCAKTD
						KAHRNVIHTSKLCFITLTPEISQ
	ļ					RSHGKSSKRVFSCFPVFLLLKH
						RRIVAQKYAELAYDLFHLGFD
						VLIIDHRGQGRSGRLLADPHLG
						HVNRFNDYVDDLAAFWQQEV
						QPGPWRKRYILAHSMGGAISTL
						FLQRHPGVCD\AIALTAPMFGIV
						IRMPSFMARQILNWAEAHPRFR
						DGYAIGTGRWRALPFAINVLTH
						SRQRYRRNLRFYADDPTIRVGG
23827	54195	В	23965	1	3357	
23828	54196	Α	23966	14682	14931	EIGPGPRPLPSPLP*ATSTSVLAA
						SGRPERTR\HAGIKIVLEDIFTL
	ļ ļ					WRQVETKVRAKIRKMKVTTKV
						NRHDKINGKRKTAKEQSPLLQE
						SLFATGDVSHNLLRALDVGLLA
						NLSALAELDISNNKISTLEEGIF
						ANLFNLSEINLSGNPFECDCGL
						AWLPRWAEEQQVRV/AAARGS
						HVCWAWLPGWPASAWHPLAG
						QWLW*GVCRLPP*QQLRHRGS
						SVLFSCPRRPA/CSQRPAAPSAS
		<u> </u>	<u> </u>	<u> </u>	l	PPARASQPSPSRA

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	:	
23829	54197	A	23967	1	714	MVKLSIVLTPRFLSHDQGQLTK
2002	"					ELQQHVKSVTCPCEYLRKVINT
						LADHRHRGTDFGGSPWLLIITV
						FLRSYKFAISLCTSYLCVSFLKTI
		l				FPSQNGHDGSTDVQQRARRSN
						RRRQEGIKIVLEDIFTLWRQVET
						KVRAKICKMKVTTKVNRHDKI
		l				NGKRKTAKEHLRKLSMKEREH
		İ			<u> </u>	GEKERQVSEAEENGKLDMKEI
						HTY/MITPSARKPLCNRVRMAA
						AEHRHSSGLPYWPYLTAETL
23830	54198	A	23968	111	216	ALLFPISTWSTPLGTQQAPQCV*
23830	134196	A	23908	111	210	PTPMPMRPVPS
23831	54199	_	23969	1	4710	FIFWFWRFVFS
23832	54200	A A	23969	3544	3723	RSHRSPPRCPPSRAPAPRRSAAA
23632	34200	A	23970	3344	3123	PPRR*PPRAPAAPRPGPP/CMAR
,						PPPARPTAGSRVPPL
22222	54201		23971	3	1119	EFGTSHRHRGTDFGGSPWLLIIT
23833	34201	Α	239/1	3	1119	1
			:			VFLRSYKFAISLCTTYLWVINTL
						ADHRHRGTDFGGSPWLLIITVF
						LRSYKFAISLCTTYLC\VSFLKTI
						FPSQNGHDGSTDVQQRARRSN
						CRRQEGIKIVL\EDIFTLWRQVE
						TKVRAKIRKMKVTTK\ATRLTK
						IKERRKTAQDHWRKLSMKERE
						HREEERQVSEAEENGKLDMKEI
						HTYMEMFQRAQALR\RRAED\Y
						YRCKITPSARKPLCNRVRMAAA
	,					EHRHSSGLPYWPYLTAETLKNR
			:			MGHQPPPPTQQHSIIDNSLSLKT
						PPECLLTPLPPSALPSADDNLKT
						PAECLLYPLPPSADDNLKTPPEC
						LLTPLPPSAPPSADDNLKTPPKC
						VCSLPFHPQRMIISRN
23834	54202	Α	23972	1	1149	

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	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ł				sequence		
23835	54203	Α	23973	93	1165	HRRSAAEAPAAAVMPFPFGKS
23033	3 1203	ľ.				HKSPADIVKNLKESMAVLEKQ
		İ				DISDKKAEKATEEVSKNLVAM
		Ì	}			KEILYGTNEKDPQTEAGAQLAQ
]						ELYNSGLLITLVADLQLIDFEGK
		1				KDVAQIFNNILRRQIGT\RTPTV
1			}			EYICTQ\QNI\LFMLLKGYESP\EI
						ALN\CGIMLRECIRHEPLGKIIL
						WSEQFYDFFRYVEMSTFDIASD
						A\FATFKGLLTRHKLLSAEFLEQ
1						`
						HYDRFFSEYEKLLHSENYVTKR
						QSLKLLGELLLDRHNFTIMTKY
						ISKPV\NLK\LMMNLLRDKSRNI
						QFEAFHVFKA\FVANPNKTQPIL
						DILLKNQAKLIEFLSKFQNDRTE
						DEQFNDEKTYLVKQIRDLKRPA
						QQEA
23836	54204	Α	23974	3	158	
23837	54205	Α	23975	3	1071	GVRASGRAVPRAVFAGMKRPC
						EETTSESDMDETIDVGSENNYS
						GQSTSS\VIRLNSPTTTSQIMAR
						KKRRGIIEKR\RRDRINNSLSELR
					!	RLVPTAFEKQGSAKLEKAEILQ
1						MTVDHLKMLQATGGKGYFD\A
						HALAMDFMSIGF\RECLTEAAR
						YLSSVEGLDSSDPLRVRLVSHL
						STCATQREAAAMTSSMAHHHH
						PLHPHHWAAAFHHLPAALLQP
						NGLHASESTPCRLSTTSEVPPAH
						GSALLTATFAHADSALRMP\ST
						GSVAPCVPLLSTSLLSLSATV\H
						AAAATATAAAHSFPLSFAGAFP
						MLPPNAAAAVAAATAISPPLSV
					,	SATSSPQQTSSGTNNKPYRPWG
						TEVGAF

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23838	54206	Α	23976	1	1416	MPGNLTRKESSSPLTDRHCHTS
						SQFSHRFNAISPFPPAELGCDHV
						ASAAADPLGVPGVATQWAGGL
		ŀ				SAALAAPRRPHSCRRLQVLGAL
						GILKSAGRVNFRCFGPLSIRGKP
				1		KTPLWEQEKPKLQTGSAGSQQ
						KRVMLKPSRGADRVLGRGSQR
		i				PAETDRETTDLRQRKRMKEGK
						ERTLRAARGETRRSNGNLARQ
						TVPEGCRSQTQALQVVRNART
		l				PPCIPNLDRPPEACSSGSTGGM
				İ		MPLRVAPSPVASGESSHLSSGR
				ĺ		GDRGLVRCKVEELRFQPVAFPE
						FIPPPTSPRADSESGDWCCTVLA
						NPSKTGSGGWEQRKTEC\PQGR
						TQNPQQRGS/RNLVLLTAGQES
						TPRRSQLTRSPAAERRTEEQGE
				İ	1	DRKNRALGCNLFPRVPRHARD
						SQRALRASTQAPKKSTFAFPGI
						HSGAVCSKRRICGVQGEPFPLPF
						ASQCSSCLKILPSPGPTKELTSY
						LLHSGCGFLDKLRV
23839	54207	Α	23977	3	139	
23840	54208	Α	23978	141	734	GSQLRRPAFGVRVALEESKGLI
						RKDSCRGRACAGANACARRRR
						PPTMSGAAWRGTPLALVRSSRT
1						RGVWAWQWSATRSPSDSCFPG
		1				VARDKEDGEPKTKHLREEEEE
		1				GEKHRELRLRNYVPEDEDLKK
						RRVPQAKPVAVEEKVKEQLEA
						A\SPEPVIEEVDLANLAPRKPDW
						DLKRDVAKKLEKLKKRTQRAI
					<u></u>	GRLDP
23841	54209	Α	23979	3	585	GTVGGVCGPRRSSSASPQEQDQ
						DRRKDWGHVELLEVLQARALF
						TAVPLSEEDRATVLKAVHKVK
					1	PTFSLPLQLPPPVNTSKLLRDVY
						AKDGRVSYPKLHLPLKTLQCLF
						EKQLHVELASRMCVVSVEKPT
						LPSKEVTHARKT\LKTLRDQWE
						KALCRELRETKNRLERQVYEG
						RFSLYPFLCLLDECEVVRMLLQ

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NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23842	54210	A	23980	1	3778	MRRRGPVGWPGAACVMSALC
						WGRGAAGLKRALRPCGRPGLP
		l				GKEGTAGGVCGPRRSSSASPQE
		1				QDQDRRKDWGHVELLEVLQA
		1				RVRQLQAESVSEVVVNRVDVA
						RLPECGSGDGSLQPPRKVQMG
						AKDATPVPCGRWAKILEKDKR
		1				TQQMRMQRLKAKLQMPFQSG
		ł				EFKALTRRLQVEPRLLSKQ\MG
						GCLEDCTRQAPESPWEEQLARL
	1					LQEAPGKLSLDVEQAPSGQHSQ
						AQLSGQQQRLLAFFKCCLLTDQ
23843	54211	A	23981	101	805	HMYTSMKPSSQPRSWISTPHPK
						ACSSLSFPLSLDNWMNINLREVI
						CTSRDGDKFWRMPECYIRGSTI
						KYLRIPDEIIDMVKEEVVAKPR
						GRGGLQQKQQKGRGMGGAG
						RVGSPLSSG\VPGAGRQPHVAS
		İ		į		LLLGVALAALPRRLPNLPGPPS
						CPLAMNRGPGRGAPRHGTQPSS
			•			QCASWRSSGHIKSPPQAPVSPM
			}			RGNGGAHLSRPLATTLGTLRGS
22211	5.010	ļ	22002	122	10.60	IQSQGRPPPPEMMNWH
23844	54212	Α	23982	133	1069	RIAWGQDRSCLQGGPFATYVP
						GLILCSLQGWYGVQAGQDLQG
						LPRSQGWARRPGGKLITCSPSP
						QDGDKFWRMPECYIRGSTIKYL RIPDEIIDMVKEE\VVAKGR\GR
						EGPCQQAESQQK/GVRGMGGA\
						GRCVFGGRGRGMGSPITG\RCH
						PDKKPGRQAGQTVSAHPDRLL
		i				RPLLARVGDFRSTVSERNLLQK
		ļ				GRGRPPRGEPPFRVLVTPSLFKV
						KAPFFLPKNRQILRVRKCLLCCF
	İ					FGFCETGPSQKLGWNPLERVLC
						LKAIFSSPFASGFSRASHFPRRG
	İ					LFLHFLGSLSEASNQTGRFLILK
23845	54213	Α	23983	1	326	PVSLECYTVPPEDNLALLQLYF
23043	134213	^	23703	1	320	RTLVTGA/LRPRWCPVLYAVTL
						WPHVNSFIFSQRPRIVDIDGKM
	1	1				GTLSCIQPFNRCLARRRTLCPNG
				1		RQETKEQKACLPGSRLFELF
23846	54214	A	23984	2	341	KATI KEAKWACTI QQKTI ETL
23840	134214	A	23704	14	741	

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23847	54215	Α	23985	114	712	DLISALMIWFHVIVIHTPYVCIYI
						PTLPLSTSFPLSQLPVSLECYTV
						PPEDNLALLQLYFRTLVTGALR
						PRWCPVLYAVAVAHVNSFIFSQ
						DPQSSVSYISIFRGRKGASN*AA
						SGPAGEEVSVKQGQWWVPAL
						GSVPPLPVLLL\DEVKAARRSM
						LQKTWLLADEGLRQHLLHYKL
į						PNSTLPEGFELYSQMPPLRQHY
23848	54216	Α	23986	357	2771	FESSCPPVGSPVGAGPTPSLYKV
						PCATAMKLLRVLASAGRNIAA
			,			RLLSSFDLRSRLCRIIAEAPQEL
}						ALPPEEAEMLSTEALRLWAVA
						ASYGQGGYLYRELYPVLMRAL
		l				QVVPRELSTHPPQPLSMQRIASL
		İ				LTLLTQLTLAAGSTPAETISDSA
						EASLSATPSLVTWTQVSGLQPL
		İ				VEPCLRQTLKLLSRPEMWRAV
		1				GPVPAACLLFLGAYYQAWSQQ
		l				PSSCPEDWLQDMERLSEELLLP
		1				LLSQPTLGSLWDSLRHCSLLCN
		l				PLSCVPALEAPPSLVSLGCSGGC
		1				PRLSLAGSASPFPFLTALLSLLN
1		1				TLAQIHKGLCGQLAAILAAPGL
		ł				QNYFLQCVAPGAAPHLTPFSA
						WALRHEYHLQYLALALAQKA
						AALQPLPATHAALYHGMALAL
						LSRLLPGSEYLTHELLLSCVFRL
						EFLPERTSGGPEAADFSDQLSL
		l				GSSRVPRCGQGT\LLAQACQDL
		l				PSIRNCYLTHCSPARASLLASQA
ł						LHRGELQRVPTLLLPMPTEPLL
İ						PTDWPFLPLIRLYHRASNTPSG\
		1				LSPTDTMG\TAM\RVLQWVLVL
		l				ESWRPQALWAVPPAARLARLM
						CVFLVDSELFRESPVQHLV\AAL
1					1	LAQLCQPQVLPNLNLDCGLRG
ļ						LTSFPDLYANFLDHYEAVSFGD
ŀ						HLFGAMVLLPLQRRFSVTLRLA
ļ						LFGEHVGALRALSLPLTQLPVS
						LECYTVPPEDNLALLQLYFRTL
23849	54217	Α	23987	291	454	KAETGSLPYTLCKNQFKMD*RL
						TC*T*NHKNPRRKPR*YHSGHR
						HGQRLHDENT
23850	54218	A	23988	1	613	
23850	54218	A	23988	1	613	

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23851	54219	Α	23989	807	1345	RCEERLLPLNLAVIALPSGLER
}						WSFFAVYNGHPGSQVAKYCCE
						HLLDHIADNQ/DFTGSAGAPSLE
						NVKNGIRTG/CPEPEVHDTERSE
						EDDHFIILACDSIWDVMGNEEL
		ŀ				CDFVRSRLEVTDDPEKVYNEIV
						DTCLHKGSQDNMCVIF\LWFPN
						APKVSPEAAKKEAELGKYRLN
						SILSDQ
23852	54220	Α	23990	344	636	SSLSLLDIWVGSKSLLL*IVPQ*T
						YVCMCLYSSMIYSPLGIYPVMG
				1	ļ	WLGQMVFLVLDPSGIATLTST
		ŀ				MVELVYSPTNSVKVFLFLRILSS
						TCCFLTF
23853	54221	Α	23991	1	2793	
23854	54222	Α	23992	18	409	SSVVEFVDPRVAPALLPQLPLES
						SSHGSVLPTMAVCLDKGHKMT
						KNMSKPRHSCRRGRLTKH\KFM
1			ļ			WDMIREVCGFAPYERHAMVLL
1						KVSKDKRALKFIKKRVGTHIRT
						KRRQEELSNVPAIMRKADAKK
						D
23855	54223	Α	23993	1	609	MHPRGPPPPAQEESHTDESPGID
						HQSLARRAGRVGRTGGFSQLL
						AQTNTSMLSSSPASCTSPSPDGE
		1				NPCKKVHWASGRRRTSSTDSES
						KSHPDSSKIPRSRRPSRLTVKYD
].	1				RGQLQRWLEMEQWVDAQVQE
						LFQDQATPSEPEIDLEALMDLST
	1					EEQKTQLEAILGNCP\RPTEAFIS
						ELLSQLKETPETQPASEISLRDH
23856	54224	Α	23994	1	204	
23857	54225	Α	23995	1	418	MWNRLCNWVTSIGWKSLEGSE
		ļ				EDRKSSQWGTLPSGAVRRGPPS
	1					SRSKNGRSNSSLHPVPGKATGT
						QLEPMRAVMGDESCKATGTEL
						LHALGAHPLHRLLKQHSHMMF
		1				QKVQVEPLRK\EPHHWQRL*PF
						SSE*NHGPSR
23858	54226	Α	23996	104	253	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		·
23859	54227	Α	23997	666	1436	WRPCFWLQAKKEAPAP\PKAK
						AKVKAL\KAKKAVLKDVHSHK
						KKKD\RKSPTFRG\PKTLCTR\KP
						PKCLQKTAPRKNKLDHCAITKF
						PLTTKSTTKKIEDNNTLVFIVDV
			:			KANKHQFKQAVKKLYDTDVA
						KNITKRWDVTSIITSCYIHLRLA
						SRLALEDLLAGLMNCLKGTFW
		l				NEESGERVETRPESRFKQQSKLI
						YTIKNPGRKKYGAEEKRAFWC
						HTDLLLHDHDEQVNHFSSPSST
		Į				FTKCKRRKPFLLREFVWIK
23860	54228	Α	23998	2	301	
23861	54229	A	23999	1	242	MGMVRKQALERTEWALVCRQ
						AWSSRMGYSTTVSPTQCQYTC
						SRGLLDSQLQLYKCTQALPEIH
						SLGHVCPSLITVKYSHYVLAAA
					:	TLFIWDNNESLRTGLLSVSTCK
						SGTRDFMRISGMVSLTSEVDNV
						EVFHYFPLHIQGKKFWAVGAI
						MTIFVISIYYEPGAHWHHMWK
						LLKVYSLESQKREEDISKRLPNF
						LKSSRRPWEDLILMLRKSPEMS
						KEYRASPLSLSAALRSKDRQSR
						HRCVTSTFHIREVRCCKSQGLP
						LCMHETAAHICP*TEGQQKGSG
						GLLYIP*TSPEIFSASESDPPMAC
23862	54230	Α	24000	2	948	CGTPTVVPGTGTSRRVRTATVE
						PTWKCRAAQQPPPPP/PPLWVK
						LPPPDTSTKPELQLEELDADSS
						WMPLHGSCPTGKLAVDARSSL
						FTEPAFPYWKKMKHTFQNILS
						WSKFTNLTYSVTQLWLLVVMA
						QTSYPHSLVGRKQVILHSSCLY
						SLDNKTSLAGDDHDDDDHDDN
						DSSPAEKGPPPITKSLDKADEQS
						HLRLAVLELGFYKPVNLLLVA
						MTGCLLQLMMKKEQCGTHGG
						EGSTLEGSRPDVPQDLFSPTLGF
						YGSTSLSADLGENVILKLYLPSF
						PLQESEPSAEHGENLWALYVGF
		<u> </u>	<u></u>			EERRSEASLAKAP
23863	54231	A	24001	265	513	
23864	54232	Α	24002	513	796	TGTAWTPPPPPLTTGAPCTPPPR
						CTARGRT/PGDSHLGGGPAATA
						GGPRTSPMSSGGPSAPGMRPPA
						SSPKRNTTSLLNSGLEPTFSFRIT
		<u>L</u>	<u> </u>			FGFM

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23865	54233	A	24003	1268	2097	TPELTGAISLSTLGDPSPFRQPSS AQMGHLLSTSLSTPPTTSNSETT QPAFATVTPSSSSVLPGLPQTSF SGMGPSAELMASTSPKQQLPQF SAAFGHQLSSHSGIPKDLQPSHS SIAPPTGFTVTGVKLQTFAVSVT VLKATCLELFVPPGGFVVSLAS GVKLQTVTPLHFSGNHPSAFYY YRLSFEMDKGKKRQFGVEKEK CGSLGGFVMLGNEEMIRRAVT LIAWTKIPFLGIREAKNPGSENI RLATILEAP\PPSWELWEQGPPG
23866	54234	A	24004	3	1163	APYYQP GEVTSWNRGRGLRIWRYFLFGP
23000	J-72J-4		27007			VQKTMRLLLSRLTYCWLEQDG RVSWRLGLEIGSLPTPVPVRLCP ANRSPPMVCGTTGRSSSNSSTE KRMCTCSSRAALHFLKGPEGG TLMEEIRALIKEPPESSLTPSTRG GHSEKAPSKNHQAATVNTTTLS RISHISHTHLLIHCTGKEQHEPY EDAATRSRPEADLSRHERLTCR HLDLGLPDMQNRQCVPGSQGL RPVGGLEGPRHWCLVRSSRLG VLPRRPQGCEDDAGGLQRQDC GQWRELVQLPTPP/VCPGGASP HPKPLPDLPDPPGSSRCFCLLPG TCPAAPPAGSPSPCSQSPGCLPP VPTPSVLECCTQVLLEDSKPQS HRNSAPHLQPGIWSHIPGSHPW CLVLAAEGSTSSV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23867	54235	Α	24005	1	2502	MEARVERAVQKRQVLFLCVFL
		1				GMSWAGAEPLRYFVAEETERG
	İ					TFLTNLAKDLGLGVGELRARG
						TRIVSDQNMQILLLSSLTGDLLL
						NEKLDREELCGPREPCVLPFQL
						LLEKPFQIFRAELWVRDINDHA
						SVFLDREISLKILESTTPGAAFLL
						ESAQDSDVGTNSLSNYTISPNA
						YFHINVHDSGEGNIYPELVLNQ
						VLDREEIPEFSLTLTALDGGSPP
						RSGTALVRILVLDVNDNAPDFV
						RSLYKVQVPENSPVGSMVVSV
						SARDLDTGSNGEIAYAFSYATE
						RILKTFQINPTSGSLHLKAQLDY
						EAIQTYTLTIQAKDGGGLSGKC
						TVVVDVTDINDNRPELLLSSLT
						SPIAENSPETVVAVFRIRDRDSG
	1					NNGKTVCSIQDDLPFILKPSVEN
	}					FYTLVTEKPLDRERNTEYNITIT
						VTDLGTPRLKTEHNITVLVSDV
						NDNAPAFTQTSYTLFVRENNSP
						ALPIGSVSATDRDSGTNAQVIY
						SLLPSQDPHLPLASLVSINADNG
						HLFALSPWTTMPLRAFEFRNNS
						PALHIGSVSTAETNIQVTYSLLP
						PRNPHLPLASLVSINTDNGHLF
						ALRSLDYEALQEFEFRVGASDR
		l				GSTALSSEALVRVLVLDANDSS
		1				LFVLFPLQNGSAPCTELMPRA\A
						AEPGYLVTKNAWLS*QLLKAT
		-				EPGLFGVWAHN/GTDRLLSERD
						TAKHRLMVLVKDNGEPPRSAT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid] - ' -
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23868	54236	Α	24006	i	1960	MEARVERAVQKRQVLFLCVFL
						GMSWAGAEPLRYFVAEETERG
		l				TFLTNLAKDLGLGVGELRARG
						TRIVSDQNMQILLLSSLTGDLLL
		İ	•			NEKLDREELCGPREPCVLPFQL
						LLEKPFQIFRAELWVRDINDHA
		l				SVFLDREISLKILESTTPGAAFLL
		İ				ESAQDSDVGTNSLSNYTISPNA
						YFHINVHDSGEGNIYPELVLNQ
						VLDREEIPEFSLTLTALDGGSPP
						RSGTALVRILVLDVNDNAPDFV
						RSLYKVQVPENSPVGSMVVSV
	:					SARDLDTGSNGEIAYAFSYATE
		}	:			RILKTFQINPTSGSLHLKAQLDY
						EAIQTYTLTIQAKDGGGLSGKC
					:	TVVVDVTDINDNRPELLLSSLT
						SPIAENSPETVVAVFRIRDRDSG
						NNGKTVCSIQDDLPFILKPSVEN
				!		FYTLVTEKPLDRERNTEYNITIT
		ŀ			•	VTDLGTPRLKTEHNITVLVSDV
		ŀ				NDNAPAFTQTSYTLFVRENNSP
						ALPIGSVSATDRDSGTNAQVIY
	2					SLLPSQDPHLPLASLVSINADNG
						HLFALSPWTTMPLRAFEFRNNS
						PALHIGSVSTAETNIQVTYSLLP
						PRNPHLPLASLVSINTDNGHLF
						ALRSLDYEALQEFEFRVGASDR
						GSTALSSEALVRVLVLDANDSS
1						LFVLFPLQNGSAPCTELMPRA\A
23869	54237	A	24007	1	966	MGHHLMIDKLVALGGLYYAIQ
						RHYATKCSVLKNDQILVIGLFM
						IQNVIYRKHFANPLSALFLQQGI
		1				ELFAAIAEIHITTVAERNHAITQI
						RLEAOTIFDSLKEWHNAIRKSL
	•					DYEALQAFEFRVGATDRGFPAL
	•					SSEALVRVLVLDANDNSPFVLY
						PLQNGSAPCTELVPRAAEPGYL
						VTKVVAVDGDSGQNAWLSYQ
ļ						LLKATEPGLFGVWAHNGEVRT
						ARLLSERDVAKHRLVVLVKDN
						GEPPRSATATLQVLLVDGFSQP
						YLPLPRAAPAQAQADSLTVYL
						VVALASVSSLFLFSVLLFVAVR\
						LCRRSRAAPVGRCSVL
L	l	<u> </u>	L		L	LCIGISIAAI YUKUSYL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23870	54238	A	24008	<u> </u>	2673	MEVMGNMGSKILITGITSTKRK
230/0	134236	A	24008	'	2073	
						VALLAKKRERKIISQRKDVTDS
						RAKRQSEEKSIGKETVVIGIGLI
			 			SKDSAAAICESRWRLSESHFER
						LIQRRMEARVERAVQKRQVLF
						LCVFLGMSWAGAEPLRYFVAE
						ETERGTFLTNLAKDLGLGVGEL
						RARGTRIVSDQNMQILLLSSLT
					:	GDLLLNEKLDREELCGPREPCV
						LPFQLLLEKPFQIFRAELWVRDI
						NDHAPVFLDREISLKILESTTPG
		İ				AAFLLESAQDSDVGTNSLSNYT
		ĺ				ISPNAYFHINVHDSGEGNIYPEL
				•		VLNQVLDREEIPEFSLTLTALDG
						GSPPRSGTALVRILVLDVNDNA
						PDFVRSLYKVQVPENSPVGSM
						VVSVSARDLDTGSNGEIAYAFS
						YATERILKTFQINPTSGSLHLKA
						QLDYEAIQTYTLTIQAKDGGGL
						SGKCTVVVDVTDINDNRPELLL
						SSLTSPIAENSPETVVAVFRIRD
						RDSGNNGKTVCSIQDDVPFILK
						PSVENFYTLVTEKPLDRERNTE
						YNITITVTDLGTPRLKTEHNITV
					:	LVSDVNDNAPAFTQTSYTLFVR
	}					ENNSPALPIGSVSATDRDSGTN
						AQVIYSLLPSQDPHLPLASLVSI
					[NADNGHLFALRSLDYEALQAF
					ļ	EFRVGATDRGSPALSSEALVRV
	<u> </u>					LVLDANDNL\PFVLYPLQNGSA
						PCTELVPRAAEPGYLVTKVVA
	•					VDGDSGQNAWLSYQLLKATEP
23871	54239	В	24009	1	993	
23872	54240	В	24010	1	4056	
23873	54241	В	24011	1	906	
23874	54242	В	24012	1	4318	
23875	54243	В	24013	41	1935	
23876	54244	Α	24014	3	328	GWQSGPVLGGLCGTAGQSCGD
						GDGIWWAQAAQAGDPAHLRL
						LTHGQPSSVPEAPVRAHQPALP
						QPRSAAARECGR/AGRSTTNSL
						RKRVASESYMIVAPPMPSSWSS
						SG
23877	54245	A	24015	114	567	VKKPHSAEWVPLWGLQPGWR
						QPFRCNSGSEKELRRPVSNSQEI
						LSPVTMRKPAVSVCCDLTLREA
						APPFQRHERTYTGEKPYKCKQC
						GKAFSGFCYTQILEKAQTGKQP
						YE/AHERTHWKQNHDCTQFGK
L	<u> </u>	<u></u>		<u> </u>		AFSFSSSLCIHEKIHTGEKLYE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23878	54246	Ā	24016	1170	491	PAQHQRGLWQSPTHQALTLML
						SVHREDWKPVLTINSIIYGLQYL
						FLESKNRWLGLERLGFCPSGLIT
						PPTCTGHRSP/NPEDPLNKEAAE
		ł				VLQNNRRLFEQNVQRSMR
23879	54247	A	24017	1	1062	
23880	54248	Α	24018	2	700	RRPTRPGGGSGGGGRMIKLFS
						LKQQKKEEESAGGTKGSSKKA
						SAAQLR\IQKDINELNLPKTCDIS
						FSRSQTTLLNFK\LVICPDEGFY
					:	KSGKFVFSFK\VGQGYPHDPPK
						VKCETMVYHPNIDLEGNV\CLN
						ILRGDWEASPLRLNSIIYGLQY\
1						LFL\EPNPEDP\LNKEAAEV\LQN
						NRR/VCFEQNVQRSMRGGLHPA
						PPTFERCLKIGLAHTHPPPRPQA
						LASPAKYLLGAMG
23881	54249	Α	24019	302	884	PMAPLSSQQPTLFPNTLTQAQQ
						VQLTKGLRKWMVQHGGDGW
	1					VVEENRTTVPGAPSQTCFVTSF/
						EVNAPAIQSWCCKKQVLDLEE
						EGLWPELLDSGRIEICVSDWWG
						ARHDSGCMYRLLVQLLDANQT
		1		1		VLDKFSAVPDPIPQWEQQCLPF
						TFTHVFFQHSRWGVRFCVFSEH
						RGPGTHSSGAGPLLSPCEQLQC
23882	54250	Α	24020	274	866	SSLLNLPSSWDYGWVSYEQAN
						CKGEQFVFEKGEYAHWDWWT
		ŀ				SSQRTDFSSLR/RPSKVDSQEPK
						V\YLYENPNFTGKKMEIIDDD\V
						PSFHA\HGYQEKVSSVWVRSG
		1				MWVG\YKYPGYRGLQYLLKKG
						DYKDSSNF\GARHPQV\QSVRCI
						HDMQGHQCGAFHPLGTQEEEE
		į				AGQKEQMEQVPFWLARGDICE
						LSGKKSVI
23883	54251	Α	24021	179	374	AASIGRSMTEKAHAGPRPGRN*
		1				ADAPSSSDTLFPPDALLSNQAG
	<u> </u>	1_	<u> </u>			RLACWLFSYGPQTKNDFHIFK
23884	54252	Α	24022	1	141	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
-				sequence		
23885	54253	Α	24023	1	506	PHSPLWKTLCSPSSGVCCGFVT
						LFCNDRFACLFQLTPPPSPAWM
		ŀ				TAAASLLASLLPALPPLNLQDA
						KFVEERRKQLQNYLRTVMNKV
						IQMVPEFAASPKKETLIQLMPFF
						VMGLQVGMYGLFHFGVMMDQ
						PQPSLTRADESPCDRVLSPCAA
			Ì			HRKQRDCCLSWKAEPE*FLVK*
						KTLCSPSSGVCCGFVTLFCNDR
						FACLFQLTPPPSPAWMTAAASL
						LASLLPALPPLNLQDAKFVEER
						RKQLQNYLRTVMNKVIQMVPE
				<u> </u>		FAASPKKETLIQLMPFFVMGLQ
						VGMYGLFHFGVMMDQPQPSLT
						RADESPCDRVLSPCAAHRKQR
						DCCLSWKAEPE
23886	54254	Α	24024	1	396	
23887	54255	Α	24025	1	1176	MVPGDLSQTSEDQSLSDFEISN
		l				RALINVWIPSVFLRGKAANAFH
ŀ						VYQQSIAGVSGIARKRALIPFGK
						STFSYVCITNIKLTDALPSWDYT
						CHFNLPKAQKPGEPTKCILTME
į						WLALLQGDPGSWIQKEFIVGKR
						GGKMDNILWDQRDVQPPGLRP
						HQLLHSGGGPLISFRNGGPILV
						WELRRKGASGKEIVGSRYFRES
						DEVLVWKNPKTLQKEDEEVNS
						THPMHTYPCTHCQVFCVTAEII
[1				NIYLQACIVLEMSMLPPGERPLF
						LLRLTTVLEVTVFDSQWSRPYH
		1				VSADPSAWKHHHHYHHHHHY
						QLSSSSPTITIINHPSPS/LSHASPS
						P/CHPSPSPAVHHHHQPSITITSR
						PSPSPAVHHHHQPSITIIIRPSSITI
22200	51056	ļ. —	24026	1240	2125	INHPSQASSIHHHHQPSIPFIVSH
23888	54256	A	24026 24027	1349 670	2125 972	KTDEWNIYRRYTEFRSLHHKLQ
23889	54257	Α	2402/	0/0	714	NKYPQVRAYNFPPKKAIGNKD
						AKFVEERRKQLQNYLRSVMNK
						VIQMVPPSSLPAPKKETLIQ\LM
						PLLRSTSTPPRRAL
22000	5.4259	-	24029	1	600	FLLKSISIFFKKAL
23890	54258	Α	24028	1	699	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23891	54259	Α	24029	221	864	VDCPRSFSAVVVWQGRFWLLS
						FIPGGFLGMTLFFASGMNWTVT
		İ				LLGFLAGNLEVEPSDTIENVKA
ļ		İ				KIQDKEG/IPPDQQRLIFAGKQL
						EDGGTLSDYNIQKESTLHLVLR
		l	ļ			LRGGMKIFVKTLTGKTITLEVE
		l				PSDTIENVKAKIQDEEGIPPDQQ
						RLIFAGKKLEDGRTLSEAIFLDV
}			[GLLLSCRTVSSCGAFILGWSVVI
						LIPVISTRDDTSTTK
23892	54260	A	24030	2	864	PSFRLAWPEGRLALLGELVCVP
						VVGRGMENGRILVSADRSGGHI
		l				CLKTLT\GKTIT\LEVEPSDTIEN
						VKAKIQDKEG\IPPDQQRLIFAG
		1				KQLEDG\RTLSDYNIQKEST\LH
		l				LVL\RLRG\GMQIFP*RP*PAKTI
						TLEV\EPSSHHRKF*RARFQGKE
		l				GIP\PDQQRLIF\AGKQL\EDGRT
						LSDYNIQKESTLHLVLRLRGGM
						QIFVKTLTGKTITLEVEPSDTIEN
				İ		VKAKIQDKEGIPPDQQRLIFAG
		1				KQLEDGRTLSDYNIQKE\STLHL
						VLRLRGGC*FFSHGIRSAQ
23893	54261	Α	24031	1	124	
23894	54262	Α	24032	97	296	
23895	54263	Α	24033	71	269	AAGYRPCPRHPGARRTAATRR
		İ				CHPSPAVARPAGCSGPP*CYAL
		1				SPDPARTRRGDPQTPPCRLPRA
						G
23896	54264	Α	24034	206	667	STRLGLPKCWDYRREPPRPAKS
						SFLTAQACEAQKTGSATRRKD
						CFLQPTFWLLPFLVLEKGAPSA
		1				CRGKRYVRAVWLSLSTGAILQP
						YPKLQVSNWL*CARDQEMPEG
		l				CL*QSCQTLSLLVLPPKEKKDV
·						*EISPWTPAPPEHQYTVRTFQSP
23897	54265	Α	24035	3	190	GTFKNDLILLISKSYY*APLPSD
						FSHSDTSS*KPSKSLKTYPSPPFF
						SAKTSLLQCPRNILY
23898	54266	Α	24036	197	474	SPLSHRGASELLSSLHLRSSCSS
						PKMAETPEAGFQIVHSASPQPG
				1		TAPGNPWAAGAGG*HCEPGQL
						AGPYRAASFFGQAWPPHPRAK
L						LGVSR
23899	54267	Α	24037	164	323	GQEIKTILANMVIPRLY*KYKKI
						SWAWWHAPVVPATWEAEAGE
						LLEPGRQRLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ŀ			sequence		
23900	54268	I IA	24038	582	1166	HCEPGQLAGHLGQPLSSGRLGH
23900	34200	^	24036	1362	1100	LMASHKLSNNYYRTRDGRRES
1						VPPSIIMWSQKALVSG\KPAERV
						YGDANHTVGNMCASRLGGYL
ļ						GFSIRIVVFIWYLGLSVVVNML
						VGRQCDWQGLVYGQPELLVFS
						GNMPGEVARGKHRNEKAPLSM
						RPASQCRVFCVLKGCSERSRHK
					İ	EQTPDTTIFKNCNTHCEGPWLH
23901	54269	A	24039	1214	3878	EQTIDITION AND THE EDIT WENT
23902	54270	В	24040	62	1242	
23903	54271	Λ	24041	74	408	
23904	54272	Ā	24042	790	1877	DEIVDPGSMKWSYKSQLNYKT
23701	3 12/2	'	21012			KRPSVFVTALTHRPDILILGTLE
						SGHSRNLTCSVPWACKQGTPP
						MISWIGASVSSPGPTTARSSVLT
	ļ					\LLPQSPQDHGTS\LTCQCT\LPG
						TGVTTTSTVRLDEAYPA\WNLT
i						MTVFQGEGTASTTLRNGSALSV
						LEGQSLHLVCAVDSNPPARLS
1						WTWGSLTLSPSQSSNLGVLELP
						RVHVKDEGEFTCRAQNPLGSQ
						HISLSLSLQNEYTGKMRPISGV
						MLGAFGGAGATALGFLSFCIIF
				1		VGVRSCRKKTARPTMG\VGDT
						GMEDT\NAVRGSA\SQGTPELN
						PRQKEQAPQNHGSANALANPP
						RPEEGEIQYASLSFHKARPQYP
						QEQEAIGYEYSEINIPK
23905	54273	Α	24043	183	1176	VHQLVFPRQHQRHSVFPKANSS
						WIHQLFPQ\QHSYGPSAVSAGQ
						HQLMVHSLFPQATPAHASISLV
į						FRQGTHQLHRVHQRRIPSAQHQ
						RHRLHQRRLPSAQHQRHRDSIS
						VCFPFRAHTRPHCDSIQRQHQL
						MRSIGLFSSGKTDPTVTLLKEA
						AAEHEICLCSRHPVNLRETKAH
						LGNAAHAVFPRGNQKKPEMCP
						YLPTHKLVVPTPPLIPLEAALRN
						IAHSRSIPPPKKFSPTPDTSDDYL
						LVFNSFPYSKLTTPKTGQGENL
						GSTRPIPCNGPPTRGPHNGAKY
						PGDVSHPVGVGTLWANRLLTG
_						QYEENGKSFFLLICTIATFQVNA
23906	54274	С	24044	178	342	
23907	54275	С	24045	86	283	
23908	54276	С	24046	91	177	
23909	54277	С	24047	11	169	
23910	54278	С	24048	209	673	
23911	54279	С	24049	164	208	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23912	54280	A	24050	182	255	DAESRRSWQPLPEPLELRP\RLS D
23913	54281	A	24051	146	456	LKNHKRSEKPCPTLTDDIPPQK KCKWPVLALSADITL*KSFSWL ILAQKAPPLSTL*PPLLPAREQM SFD\LIFLYLPKSYKVAPPLSSFT ESLFGLSPPAPR
23914	54282	С	24052	90	215	
23915	54283	В	24053	1	1059	
23916	54284	С	24054	129	215	
23917	54285	c	24055	19	347	
23918	54286	c	24056	228	260	
23919	54287	В	24057	406	602	
23920	54288	С	24058	7	375	
23921	54289	A	24059	322	400	FLLSRVPSV*DPTENQTVQLTW EWLP
23922	54290	A	24060	818	1078	ILDVFSYTMKIEPSPLRSYPISPH YNL*QLPP*LDP*ESGYKTPLSA LLLTFLLYISSFASHKVSSSSVD PLPTCAYLPIGQAHVH
23923	54291	С	24061	1	411	
23924	54292	Α	24062	1	861	
23925	54293	Α	24063	425	484	
23926	54294	A	24064	1844	2097	HSAGTKLRLVNFLLTLRRLVNS VMRLKKSFMMVQRMVSIPKV WSISLGFLLSFVPSVWDPT/ENR TVQLT*HPLPEPLELGPRLSD
23927	54295	A	24065	729	807	LFSCSCPPKPPRP*LTAEKGGLC IFL
23928	54296	В	24066	1	1317	
23929	54297	В	24067	509	810	
23930	54298	Α	24068	-	385	MDLPSTVRVTRSSASVMVQGG SEAIRQRQSSAAKPRRSGKESKS LGPEFQGLWEWLPEIYFSQFRR LGNPKSRHQQIQCLMGACFVV TERLSFQDSLDLAEPRHSYPLQ VKRLDLPTVLETETDLEKLDPF TTIPSPVETLQCLFVSLRIKSRVP *FWISQPPEL*EINFWQPLPEPLE LWPKAL
23931	54299	В	24069	1	774	
23932	54300	Α	24070	350	640	SLQGCLSDNSPTFQRCQTTQGR LPWSFTLSGKSHFSGGVARACY TCQKSGHWA\RNASSPGFLLSR VPSVSDPTENRTVQLTWQPLPE PLELWPKAL
23933	54301	В	24071	85	331	
23934	54302	В	24072	1	1844	
23935	54303	В	24073	1	1169	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	defendit, (=possible interedide insertion)
23936	54304	Α	24074	63	220	NLTGNIWNREWYSHFDFDLLD
						PVGFSGSIAKMPKRKLFKCHSV
22027	54205	 	24075	407	909	*GHWRSKSE
23937	54305	В	24075	497	898 678	CCCECDEDETIALODCOONAL
23938	54306	Α	24076	3	078	GGGFSDRDRTIALQPGQQIWM WGIGKQVGGCGWLRGKNRVV
						GAAKDQSGTGEKFYFIPKAMG
				ļ.		AMEGSGQRRTLPDPGAHRNPS
		i				GGYSATDCEGSRPAPGEINSHA
						SHTKPVWWFFTQTRMKFGAVT
		i				RIGGLPWEVNPLSSCSLLRE/KD
		1				PPTTSGPQSNQPRKH/LTNFKSG
		l				A/CYTCRKSGHWAKECPLPGFL
		1				LSRVPSVWDPTEN/RTVQLTWQ
]				PLPEPLELWPRLSD
23939	54307	В	24077	522	873	
23940	54308	В	24078	329	530	
23941	54309	С	24079	147	439	
23942	54310	Α	24080	3	3889	
23943	54311	Α	24081	2	394	
23944	54312	Α	24082	1	197	
23945	54313	Α	24083	1 .	205	CLRAGPHALYLLC/NIPEFTFL\S
		1				RADF\ARSQLVDLLTDP\FQ\QEL
23946	54314	A	24084	134	182	EELLQVGAVTLILRILPCLPCLP SCFIPSSQPVAGTERSRFLGPQY
23940	34314	A	24084	134	162	FQTPHNPSGPGLGNQLMR*AF
23947	54315	A	24085	257	481	POTTINI SOF GEORGEWIK AT
23948	54316	A	24086	415	630	
23949	54317	A	24087	123	340	
23950	54318	В	24088	49	270	
23951	54319	A	24089	1	900	RSRVRREVVALRRGLACIPSLG
						VWRFPSVGVLAVSDRWCLAVS
		1				DRWCLARATLSCFGCAMPMYQ
1						TRRQENDLRTASIAV*RRKQQN
	ı	İ				DH*KQRRWQNIQRKGPKRYIVI
		l				AGNSQSHQPMIFSMLRKLPKVT
		l				CRDVLPEIRAICIEEIGCWMQSY
					1	STSFLTDSYLKYIGWTLHDKHR
						EVRVKCVKALKGLYGNRDLTA
					1	RLELFTGRFKDWMVSMIVDRE
	-				1	YSVAVEAVRLLILILKNMEGVL
	:					MDVDCESVYPIV*ASN*GLASA
						VGEFLYWKLFYPECEIRTMGGR
	<u> </u>		<u> </u>	İ	<u></u>	EQRQSPGAQRTFFQLLL

23952 54320 A 24090 I 812 MARTPGA VHRHSRK AKSDMQH LYGNRDL M\VSMIMI LILKNMEG LFYPECEII QRTFFQLI YLVDNLW TSLLLEKE LLGESPGI PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLTHISI WEQRNLG VEFEGTRI VDTEIP/ES PCPPFPRR KHPPFSPL QMIVGGR \SELASFLM CSDSIPGA	ELAVAKYPKKGSQA QSEPPANDIFNAAKA IREVRVKCVKALKG TARLELFTGRFKDW DREYSVAVEAVRLLI GVLMDVDCESVYPIV RTMGGREQRQSPGA JLSFFVESKLHDHAA /DCAGTQLKDWEGL
VHRHSRK AKSDMQH LYGNRDL M\VSMIMI LILKNMEG LFYPECEII QRTFFQLI YLVDNLW TSLLLEKE LLGESPGII PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	QSEPPANDIFNAAKA IREVRVKCVKALKG TARLELFTGRFKDW DREYSVAVEAVRLLI GVLMDVDCESVYPIV RTMGGREQRQSPGA LLSFFVESKLHDHAA 'DCAGTQLKDWEGL
AKSDMQH LYGNRDL M\VSMIMI LILKNMEG LFYPECEII QRTFFQLI YLVDNLW TSLLLEKE LLGESPGI: PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRI VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	REVRVKCVKALKG TARLELFTGRFKDW DREYSVAVEAVRLLI GVLMDVDCESVYPIV RTMGGREQRQSPGA JLSFFVESKLHDHAA JDCAGTQLKDWEGL
LYGNRDL M\VSMIMI LILKNMEG LFYPECEII QRTFFQLI YLVDNLW TSLLLEKE LLGESPGI: PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLTHISI WEQRNLC VEFEGTRI VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	TARLELFTGRFKDW DREYSVAVEAVRLLI GVLMDVDCESVYPIV RTMGGREQRQSPGA JLSFFVESKLHDHAA DCAGTQLKDWEGL
MVSMIMI LILKNMEG LFYPECEII QRTFFQLL YLVDNLW TSLLLEKE LLGESPGI: PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLTVHISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	DREYSVAVEAVRLLI GVLMDVDCESVYPIV RTMGGREQRQSPGA JLSFFVESKLHDHAA /DCAGTQLKDWEGL
LILKNMEG LFYPECEII QRTFFQLL YLVDNLW TSLLEKE LLGESPGII PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	GVLMDVDCESVYPIV RTMGGREQRQSPGA LSFFVESKLHDHAA 'DCAGTQLKDWEGL
LFYPECEII QRTFFQLI YLVDNLW TSLLLEKE LLGESPGII PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	RTMGGREQRQSPGA LSFFVESKLHDHAA DCAGTQLKDWEGL
QRTFFQLI YLVDNLW TSLLLEKE LLGESPGI: PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HIS] WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	LSFFVESKLHDHAA DCAGTQLKDWEGL
YLVDNLW TSLLLEKE LLGESPGII PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRE KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	DCAGTQLKDWEGL
TSLLLEKE LLGESPGI: PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	·
LLGESPGII PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	
PPPKRIGSI QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	QSTCHMEPGPGTFH
QCWFPVE 23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	SAIVQGAIEKSGERSS
23953 54321 A 24091 152 1209 MPLKNVQ LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	MKEGPMVLPLALDT
LIFSPLHW HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	
HFSNPLLL WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLN CSDSIPGA	NITHFPLEKELALPL
WTLT\HISI WEQRNLC VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	CVSGYLSLPLPNPKP
WEQRNLC VEFEGTRM VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	QVILPALTLVYFSIL
VEFEGTRN VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	KSDASPGECGS*MG*
VDTEIP/ES PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	PSILPLQSPLCPTEAA
PCPPFPRR KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	//VAFCELC\QSCLSD
KHPFFSPL QMIVGGR \SELASFLM CSDSIPGA	R*VLAQSGRKKAEY
QMIVGGR \SELASFLM CSDSIPGA	N*QCHYLPWETFCPF
\SELASFLM CSDSIPGA	QAFVLLSDLLLIFSP
CSDSIPGA	DFLRPLVFFPE\ATLQ
1 1 1 1 1	MDHVFIQPGRPGQW
	Q*GLGLGCSSKI*GR
1 1 1 1 1 1	CLPCEPGDSQEDHLQ
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	RLLAGFCKLLLYGV DVFKHYNKVHQGPS
1 1 1 1 1	MPRFSAATV
	HFPPEECPPPIPTAH
1 1 1 1 1 1	TWELFLQQLQEVVV
1 1 1 1 1 1 1	LEAGAHAL\YLLCNP
1 1 1 1 1 1	DFARSQL/V*DLLTD\
1 1 1 1 1 1 1	ELLQGRELGAGTMG
	GGGCKTGGMWDL
1 1 1 1	YCGIFFSFSILPSQIED
1 1 1 1	HSETPLCLLQVSGFP
1 1 1 1 1	GVCGVGSLCSFPV
	REMSAPSRSLGRFL
1 1 1 1 1	PPSLALVPPCRCTRL
KTYHGRG	APLRVELPTCMYRL
SNVQGRS	GGPAPGVGHLQSLV
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	KQDQ\DAGFL\ELVN
FFI*S\CGM	*RNSGDYPLTAPGL
SWKKFQG	SFCEFVGTLVCRCQ
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	PMDNLISLLTGFSDS
QVCAF\RH	TSTLAAMKLMTSLV
	HQDINQRQYEAER
1 1 1 1 1 1	APERLESLLEKHKEL
H*YQAKN	
23956 54324 B 24094 211 3841	LSRARCASTCGASH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or pepulae sequence	determine, it possesses adelectine insertion,
23957	54325	Α	24095	1	1798	MSLLAFKIILVGFEGSKARVQT
						DLFYRSLVLFLLLQICFPLSCFG
						NWKGEEGGMQRWVLGKNEKI
1						ASPHRLPAPTARFAVSLQLNLRI
						LGAFQFGPVLFVLWRPLDRDN
		1				ASPQSEFLNSCGITAYLVGKRE
		1				GGVERGGGGRSSERCRTRLSFS
						PRHSHVFSTSRFCKILKKSPGAN
		1				GGRCRARGGGLHGARAANPPT
		1				PIAEPARLPACLSRAEPEPQAAE
		1				PAPPPVGVVRSRPLSHRRHPVV
		1				CPAAPIRLINKRLAHSARPAGIC
[CVSRSGLSALAAAGTASMFQLP
						ILNFSPQQVAGVCETLEESGDV
						ERLGRFLWSLPVAPAACEALN
						KNESVLRARAIVAFHGGNYREL
						YHILENHKFTKESHAKLQALW
}						LEAHYQEAEKLRGRPLGPVDK
						YRVRKKFPLPRTIWDGEQKTHC
						FKERTRHLLREWYLQDPYPNPS
						KKRELAQATGLTPTQVGNWFK
						NRRQRDRAAAAKNRSQTSPWA
						EAFRFPECGSRLAKHFAEEPGLI
						PQAQMLLTILFAFRSGRPDPTRS
						REHLLRRKPQADAPLEM\LQQQ
}						VLSQGSGRALRAEGDGTPEVL
		L				GVATSPAASLSSKAATSAISITS
23958	54326	Α	24096	3	1034	GLHGARAANPPTPIAEPARLPA
						CLSRAEPEPQAAEPAPPPVGVV
						RSRPLSHRRHPVVCPAAPIRLIN
			i			KRLAHSARPAGICCVSRSGLSA
						LAAAGTASMFQLPILNFSPQQV
		ł				AGVCETLEESGDVERLGRFLWS
						LPVAPAACEALNKNESVLRAR
						AIVAFHGGNYRELYHILENHKF
						TKESHAKLQALWLEAHYQEAE
				ļ		KLRGRPLGPVDKYRVRKKFPLP
						RTIWDGEQKTHCFKERTRHLLR
						EWYLQDPYPNPSKKRELAQAT
				ł		GLTPTQVGNWFKNRRQRDRAA
						AAK\NRLQQQVLSQGSGRALR
				†		AEGDGTPEVLGVATSPAASLSS
L	<u> </u>		l			KAATSAISITSSDSECDI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23959	54327	A	24097	3	1024	GTRVGVAWRSVRLLLGPGAGL
	,					RGGVVLKVSPSPCRGGRRVPVR
						AEGARGRGPINKVPGASSGAA
						VGAARLTAPLLAKAMASKLLR
						AVIL\GPPGLGKGAPVCQRIAQN
						FGLQ\HLS\SGHFLRENIKA\STE
						VGEMAKQVYREKVFLVPDHVI
						TRLMMSELENRRGQHWLL\DG
ļ				ļ		FPRTFGQAEALDK\ICEVDLVIR
						LNISFETLKDGLN\RRWIHP\PSG
[\RVYNL\DFNPPHVHGIDDVTGE
						i .
1						PLVQQEDDKP\EAVAA\RLRQY
						KDVAKPVIELYKS\RGVLHQFS
		1	:			GTET\NKIWPYVYTLFSNKITPI
						QSKESILTLAQWEEPGRCGSFIP
22262	54300	ļ.—	24000		150	IVWCSNWCCCPNLEA
23960	54328	A	24098	2	178	DG A L ED LOUD I DUED IDLOGGOVI
23961	54329	Α	24099	96	340	FGALTRIG\DLPWEINPLSSCSLL
						PEKDPPMTSGPQTDQPKEHLTN
						FKSAARRSSRTSSHRNLLHVLEI
00000		ļ	24100	_		WQLDQGMPAARDS
23962	54330	A	24100	3	155	
23963	54331	A	24101	1	162	
23964	54332	A A	24102 24103	167	1032 476	TAMIL TOSI ECCI ETETIMANEC
23965	54333	A	24103	107	476	TAMLLTQSLFGGLFTPTHMKFG
						AVTQIG\DLPWEINPLSSCSLLR
						EKDPPTTSGPQTHQPKEHLTNF
						KSDHHGCRASGNSHSGRMLLIF
23966	54334	A	24104	371	772	SPSRGLYACSCVKT
23967	54335	A	24105	146	312	QKRSIPNSSPIP*SHSLLANCTK/
23701	3 1333	1	21103	"	312	DHHFLNGSKSFQPQLLSCKFS*
	ļ					GQHLFQGALCT
23968	54336	A	24106	76	441	ogne. goneon
23969	54337	A	24107	1	1284	
23970	54338	A	24108	150	371	EESRAAGIPWPSGQISGTCSKL\
						CGRRFWRNAWPLRFRHLEVLV
						CWRCGCGLSHSGGKELQLFSII
						VHLEGEVN
23971	54339	A	24109	280	351	
23972	54340	A	24110	1	483	
23973	54341	Α	24111	129	363	FGVVTQIG\DLPWEINPLSSCSL
						LCEKDPPTTSGPTKKHLTNFKS
						SKRPLFTLFSNLPHYPSTSFSFQS
L						WRYTSISPFS
23974	54342	A	24112	2	111	
23975	54343	Α	24113	3	808	· ————————————————————————————————————

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23976	54344	A	24114	497	815	TALLLTQILFGGLFIWTRVKFG ALTWIG\DLPWEINPLSSCSLLH EKDPPTTSGPQTDQPKKHLTNF KSACFERIKACYHSPATAWPFK AYKLSLQFPHFTCPKTR
23977	54345	A	24115	l	753	
23978	54346	Α	24116	1	513	
23979	54347	A	24117	625	701	IHVAHKLAVVMPGCKRGCLSL FYFQLIWSHKGWVRSGEASHR EMAGLLLLCSS*PPSPCETLPRS EPSLCDSRSAESRTKTSNLSCIP ASLQQVCGPREFIWSGNHCHPF LPITRCYGCLCYHYCMENSIL
23980	54348	A	24118	458	605	VVEEGIEDKRVYRLGTTGWIGK TIWLVRRRS*TNL*DLSGFWTG KMGEL
23981	54349	А	24119	566	769	IKEKGGLFSFSASNLWKQVTSS FLCSPLPLSLLKSSKLLANRDKY RMCGPVPANGNRTQQ*SKHVR
23982	54350	Α	24120	345	664	GEIAASQEPAGGFSGSSAQPCPL PWTGPPTTQAPGH/CNSAPSSRD NASSNGGRRQRVPTVPGGGLIL QREHILLPLGPALRAAGKEPAT AATIPESPGGDSDDSRT

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23983	54351	A	24121	2	2204	IDMIFTPGPPSTPKHKKSQKGSA
25705	.55.			_		FTFPSQQSPRNEPYVARPSTSEI
						EDQSMMGKFVKVERQVQDMG
						KKLDFLVDMHMQHMERLQVQ
		ŀ				VTEYYPTKGTSSPAEAEKKEDN
						RYSDLKTHCNYSETGPPEPPYSF
		ŀ				HQVTIDKVSPYGFFAHDPVNLP
						RGGPSSGKVQATPPSSATTYVE
						RPTVLPILTLLDSRVSCHSQADL
1]	QGPYSDRISPRQRRSITRDSDTP
					-	
						LSLMSVNHEELERSPSGFSISQD
						RDDYVFGPNGGSSWMREKRYL
						AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILQKQVG
						QSITSMLGFLSRGPSMKLCMGL
						ACVLSLWNTVSGIKGEAKKEK
						GMTFLPTTDSKKFFSLLSVTSYS
1		1	1			SFAFHKFSVAVYNISNLKTVDP
			Ì			AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
						SVNQSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
						SGVLALLLTQSLFGGLFTRTRM
						KFGAVTRIG\DLPWEINP/LSSCS
						LLHEKDPPTTSGPQTDQPKKHL
						TNFKSAARPTFLGQGQVPLNPF
						SFTLSGKSCFPRRQEPPNRLFPH
						PNLLSLCPNPLFLCPNPFSTFLE
		:			:	GKNHHPHPFSVSQVFSLGLPPS
						LWEHSPMAHRLRVQPRQQPPDI
		İ	Į			HKQVISLQSGHSLLVKSQSWSQ
						DPIKFQLPHHHKNTASVCINPSV
23984	54352	Α	24122	2102	2529	ETTMSFIFSLLYPFRMHLSFLIVS
						PALRSISRRALAAFPRGRSLDLQ
]	PAMPEPPTPS/CGLLCVPS\PPMS
						AAPCSKVPQSHLFAFMSCNTAK
						VCSFTAEASETTSPPGGTNNSR
						RATLRAVTLTRRSATLTRRSAA
						SHRSSTV
23985	54353	Α	24123	299	567	TAMLLTQSLFGGLFTQTRMKF
				1		GAVTQI\ADLPWEINPLSSCSLL
					1	HEKDPPTSSGPQTDQPKEHLTN
						FKSGETKEMHFIRGPKTPAPVM
23986	54354	A	24124	684	822	KVKRQSYSQNTFYYPICSRH*IK
23900	37337	``	127127			LQKLNSGPQTPQQDLINLAIKV
		<u> </u>		l	<u> </u>	Indiana du donnument

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23987	54355	A	24125	1163	1826	TVLLLTQSLFGGLFIWTRMKFG
23767	34333	``	24123	1105	1020	AVTR/IRGDLPWEINPLSSCSLL
						HEKDPPTTSGPQTNQPKELLTN
]			FKSVPTPSVAESFQSSFSTDPSD
			1			PSSPPQAAPRQAKPGPNSSSASA
				1		PRPYNPFITSPPHTWSGLQFHSA
ļ						TSPPPPAQQFPLKKVAQVKGTV
				1		KWLKTDAARSPQKPTGTITDAL
						DNSHSGGPIMDRPLLVKSPKQF
						LRLLVFSKTFISLTILNLQE
23988	54356	Ā	24126	636	922	TATLLTQSLFGGLFTRTRMKFG
23900	34330	^	24120	030	922	AVTRIG\DLPWEINPLSSCSLLRE
						KDPPTTSGPQTHQPKEHLTNFK
						SDKGDMFYPWTQNSSAGHGLG
						1
22000	54257	 	24127	728	1013	RQPSLGV MKFGAVTRI/RGDLPWEINPLSS
23989	54357	Α	24127	1/28	1013	1
						CVLC/CREKDPPTTSGPQTDQPK
1						EHLTNFKSGKRPAFYSLLQPPSL
ŀ		ŀ				SLNLFLLSILAPHFNLSLLLISIPF
22222	540.50	 -	0.4100	1.	2002	IFW
23990	54358	В	24128	1	3993	
23991	54359	В	24129	105	420	
23992	54360	В	24130	45	2866	
23993	54361	В	24131	1	1680	
23994	54362	A	24132	1257	1403	
23995	54363	A	24133	1	2130	
23996	54364	Α	24134	1	725	RAADGVITLEDNDPCVKKDSSR
23997	54365	Α	24135	2825	4123	1
1		}				FPYEDRLDLVLKGTADIPRLTV
						HRGSEYIISRATFPCYFIKEQSVI
						NHCYTEIDLKIFRQYLAPALGV
						THRFVGTEPFCRVTAQYNQDM
l						RYWLETPTISAPPIELVEIERLRY
						QEMPIYASRVMFIFRPDNTALP
						RNDTPQMVKLFMDRFQVVKDI
						GVIELKVVEDQRTRAVMIKFRA
						FVKEGAVIVIRFDNKEVAFAQM
						SGNLTVPRSTSDYKARDIFFQY
-				1		RFLTRISTGHGVANNHQIRLRL
						ELAGIVPLNQLNALLLKVYPLS
		1				KLCNSPKKCQLESVFRLTQWV
1						VKKPISNRQSGLGGIVLPGLFRV
}		1				LSPGNLSPIKGTVILPNEVQPGF
1		1				LGNLVGPI/RGWIGLS*IYLPHGF
1						TG*QNSAGLADH*RASASPERA
						GVYPIHNERCAHRASRYY*KW
						HPDSVAADHLLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23998	54366	Α	24136	137	1199	ISSGGPRELGLKMIIVPGSQLSQ
						LPPGLDSTEFAAIVGNLLDNAF
						EASLRSDEGNKIVELFLSDEGD
						DVVIEVADQGCGVPESLRDKIF
			}			EQGVSTRADEPGEHGIGLYLIA
						SYVTRCGGGITLEDNDPCAPLT
						LLIVEDETPLAEMHAEYIRHIPG
						FSQILLAGNLAQARMMIERFKP
						GLILLDNYLPDGRGINLLHELV
						QAHYPGDVVFTTAASDMETVS
						EAVRCGVFDYLIKPIAYERLGQ
		1				TLTGFP\QRKHMLKSIESATQKQ
İ						IDEMFNAYARGEPKDELPTGID
						PLTLNAVRKLFKEPGVQHTAET
						VAQALTISRTTARRYLEYCASR
						HLIIAEIVHGKVGRPQRIYHSG
23999	54367	A	24137	2	243	RPGRGLGQRGIWEPAA\LLDSV
						AQQQGAWLPHVADFRWRVDV
						AISTSALARSLQPSVLMQLKLS
						DGSADLEKRCERRLQD
24000	54368	Α	24138	1	1483	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFPPVI
			ļ :			GVAAHKGGVYKTSVSVHLAQ
						DLALKGLRVLLVEGNDPQGTA
						SMYHGWVPDLHIHAEDTLLPF
						YLGEKDDVTYAIKPTCWPGLDI
						IPSCLALHRIETELMGKFDEGKL
						PTDPHLMLRLAIETVAHDYDVI
						VIDSAPNLGIGTINVVCAADVLI
						VPTPAELFDYTSALQFFDMLRD
						LLKNVDLKGFEPDSDLRYLQSC
						EWGDTGIFLKWSCIPAEKGLQT
		1				SGAWVEAADIPLLPGSPRRLSP
					1	QAGSRGGQGPKHGQQCLKMP
	}					GPRAPGLQGGSNRDP/AP/GPAV
					1	ERAPAAALSSTTIWMPMSLCP
						WRPVLAACTSMTVRGRSTMYL
						PTTTGNAGCTWPKASLATRWL
						SSPMGRQARSLMLGAQVTLSW
						DRSMPWRRRGRSSGRNLSTI
		ــــــــــــــــــــــــــــــــــــــ	L	<u> </u>	<u> </u>	DIGHT WRIGHTHOOD GRAED IT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24001	54369	Α	24139	727	1498	GICIWDRPPGLADQRRSHSHGW
						WGELATPLPWHH/HLSDSHSG\
						RVSFLGAQLPPEVAAMARLLG
						DLDRSTFRKLL\KFVVSSLQGED
						CREAAAAVLGSAPTCRR\AAGC
						\LLAGMHTLLQQALRLPPTSLK
						PDTFRDPAPGALHPPRPGRGL\G
						QRGIWE\PSGPSFDSVAQQARV
		1	1			AWLPHVA\DFR\WRVDVTIST\S
		1				AL\ARSLQ\PSVLM\QLKAFQNG
						SSIPAFEVPPQPKF\QEL\RYSVA\
						LV\LKEMADLEKK\CE\RNLQD
24002	54370	A	24140	1	312	
24003	54371	A	24141	2	524	
24004	54372	A	24142	144	494	AGSCPFAAGPGPLGSGRCFVLIR
	[-	2			FILI\QNRAGKTRLAKW\YMQFD
						DDEKQKLI\EEVHAVVTVRDAK
						HTNFVGVSGTFKIILPPLCWPLL
	1					LASVWMFNDNKPGLYLGGPFH
						NFRGRS
24005	54373	Α	24143	1	428	
24006	54374	A	24144	108	422	
24007	54375	A	24145	3	387	
24008	54376	A	24146	3	337	TRLMLLDRSGQGKVYGLVGR*
		1		-		RFQQMDVLEGLNLLITI*RNRN
				1		KLRVYYLTWLRIKIMDNDPKLE
						KKHDWTTAGYMEGCGHYRVG
						KYERIKFLVIALKSSAEVYAWA
						PKPY
24009	54377	A	24147	1	340	ERTRMNKQHNSTLAKSRPGNT
[15 15 //	_	I	1		GPEPLMPQASPGPPGPLSQTPP
						MHRPVEPQEGPHKSLVAHRVP
					1	LKPYAAPVPRS\QSLQDQPTRN
						LAAFPASHDPDPAIPAPTATPSA
		İ				RGAV
24010	54378	A	24148	335	869	ATPTLILSQKSHFWPQPLSSVLK
7010	37376	1'	27170			LTSGPEISLQSSPSPICRMKKRR
				1		LSPLASGLES*VKSGEWCEVLS
	1					RAPLFP/HNPSQAAP/EWEMQG
						TRPSIPHPLGLAGLPTQLQQGRL
		1				WGYRNPASPFRTRVAARDCAG
						KQVRTGPDARLEGSRPGVRAG
					1	LGWGRGYSERVQTCAKSSVSPS
24011	54379	В	24149	158	5640	LUWUKU I SEK VŲ I CAKSS VSPS
24011		-				
24012	54380	Α	24150	1	1854	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24013	54381	Α	24151	1	1510	MPCGKQGNLQVPGSKVLPGLG
						EGCKEMWLRKVIYGGEVWGK
						SPEPEFPSLVNLCQSWKINNLM
						STVHSDEAGMLSYFLFEELMRC
						DKDSMPDGNLSEEEKLFLSYFP
						LHKFELEQNIKELNTLADQVDT
						THELLTKTSLVASSSGAVSGVM
						NILGLALAPVTAGGSLMLSATG
						TGLGAAAAITNIVTNVLENRSN
						SAARDKASRLGPLTTSHEAFGG
		1				INWSEIEAAGFCVNKWVKAIQG
				1		IKDLHAYQMAKSNSGFMAMV
						KNFVAKRHIPFWTARGVQRAF
						EGTTLAMTNGAWVMGAAGAG
			:			FLLMKDMSSFLQSWKHLE\DG
					•	ARTETAEELRALAKE\LEQELD
						RLTQHHRHLPQKASQTCSSSRG
						RAVRGSRVVKPEGCLATSQPH
						QHQQERGDRPREDTDNEKRGG
		1				EKWSDFGSVLKGEPTGFVHGL
						DLWCEGKKSQDNGGLWPEQL
						AGWSCWHLKWTGGSRSGLPVP
						VLDAACTGTLSPDLTSWRYHPS
						SIPLPKQPLQHGF
24014	54382	Α	24152	3	515	VGTQAGAAPGPVQFLSSAWVA
		1				L\HETELPPLTAREAKQIRRE\IS
		l				R\SSKWVDMLGDWEKYKSSRK
		1				LIDRAYKGMPMNIRGPMWSVL
		1				LNTEEMKLKNPGR\YQVRSARA
		ŀ				QPTGQAVSGAQVSSWRERQDH
						PGELGVKIMKEKGKRSSEHIQQ
						MDLDVSGTLRRHIFFRDRYGT
24015	54383	Α	24153	1	1317	
24016	54384	Α	24154	1640	4841	TITGSSPLAGGHHSTCRGRMDM
			1			VENADSLQAQERKDILMKYDK
		1				GHRAGLPEDKGPEPVGINSSIDR
		1	1			FGILHETELPPVTAREAKKIRRE
		1				MTRTSKWMEMLGEWETYKHS
						SKLIDRVYKGIPMNIRGPVWSV
					1	LLNIQEIKLKNPGRYQIMKERG
	l	i				KRSSEHIHHIDLDVRTTLRNHV
		i i	1			
, ,	•			:		FFRDRYGAKQRELFYILLAYSE
						FFRDRYGAKQRELFYILLAYSE YNPEVGYCRDLSHITALFLLYL
						1
			_			YNPEVGYCRDLSHITALFLLYL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24018	54386	A	24156	479	3902	GQVDEDVSVKMDTVPGVNLSC
24010	34360	^	24130	479	3902	ILNEMRDQDKKLVEKSYKDAK
						•
						GCFFSMGEPSLIDPCLEVCYGT
					:	QLAQLQGLIRSMEQQLCELCCD
						AEHQDHEHQVLLDVKTRLEQEI
						ATYSRLLEVEDAQQGPIVMRA
						MTSECHWCLAPIRSPDCVCVLR
						QLHTRSGIHKRFCGLKGFLPER
						HNPGQLIHQNQSGCCSRMYPFD
						RKLGISVALNIPGGRSYRDLWC
						LTAVHIHIPGIDHHRLFTLGRM
						DMVENADSLQAQERKD
24019	54387	Α	24157	780	983	QHQSLIGHKYQRSHRSL*AEPE*
		Ì				LECHQPDLAEPADGPPLTTRYG
						IVNAEMGLEQLLPKGPSLKEAK
24020	54388	Α	24158	773	2016	DWPTLRENRKEPTLKYWGLVP
	'					LISGMPTWFFFFLSACGNLIPSG
		1				GIRVPMENML/RVFRAVEEHCP
		l				WFPEKGTLSVEQWDSVGAKFQ
						ELVPTGNYVPITVWGDWALLL
						QFSESGDPLPLPQLSSPTGPSLS
-						DQPLPSPTPPPPDDVENSISNSG
						DFGLTLPPGDLILFPEEPLLAAS
						AAPNRTALGHIYANSSLFKPLQ
		}				HLPLESANGSGAKLQFTYNSAG
		ļ				PSPSSAAPRPPVVSVPQPVTLPS
						TQAASLYPSSHMDTSNHQCTSA
						SSAPPMPLSHTLIPSDETTCNVF
						NYPLKKPTGRELREAKNKGSQP
						KVHEKTNPGNIRCISTEERPHEN
						KGFNHLLEWLAELKTTIHLLDC
						WFIAKDVNREQPDGRDAGDKR
						LCVKFLREIAANTYKTEVGAAK
						VMEILLHSHGATTTIAN
24021	54389	Α	24159	365	675	
24022	54390	Α	24160	1	1191	

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SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or peptide sequence	deterion, (=possible nucleotide insertion)
24023	54391	Α	24161	363	1434	QQGVCSGEKGGHWAPRQVGV
						YLLPGRVGCVSSRVSPSFPGDG
						LDSGLARRGSAVSALASGLVEE
						PMLGPPFHPTPRFKAVSAKSKE
						DLVSQGFTEFTIEDFHNTFMDLI
		}				EQVEKQTSVADMLASFNDQST
						SDYLVVYLRLLTSG\YLQRESQ
						VLRALHRRVDGTVQRSSAKQE
						VEPMCKESDHHSHHCAGPGPS
						ACPSRVGVHGTAARAAPPIRTS
						F/H*GVPSPKVYLLYRPG\HYDI
						LYKLGLGSSPLGCPGCPLLARA
						LGHCYRGFSVVVNGPISPPSSCH
						MTPPPCFIKGGAGGEPCVRVPA
						LLPAWLLCLLPPPPRWVPLLFT
						YLLLSFPNRSRFEGPGLLEAPPA
						SLGSASFPS
24024	54392	Α	24162	2	888	VFRSRHQGCLKMAAEEPQQQN
						QYPLGSDSEGVNCLAYDEAIM
						AQQDRIQQEIAVQNPLVSERLE
						LSVLYKEYAEDDNIYQQKIQ\D
						LHKKYFVHPQDQGLTGTCFYS
						GFRDSPTWEGTCWIDSKELQRF
						QGLVS/SQRSKGKTLVSQGLSL
			İ			EFHNFEGFSTTTFI/VDLIEQVEK
						ADLCRRPCLASFNDQSTSD\YL
						VVYLRLLT/SRAYLQRESKFFEH
1						FI\EGGRTVKE\FCQQE\VEPMW
						QGEATTSTSLALAQALSVSIQV
}						EYMDRGEGGTTNPHIFPEGSEP
						KVYLLYRPGHYDILYK
24025	54393	Α	24163	3	819	EGATMAAPPQPVTHLIFDMDG
						LLLDTERLYSVVFQDI\CHRYD
						KKYSWDVKSLVMGKKALEAA
						QIIIDVLQLPMSKEELVEESQTK
						LKEVFPTAALMPGAEKLIIHLR
				-		KHGIPFALATSSGSASFDMKTS
						RHKEFFSLFSHIVLGDDPEVQH
						GKPDPDIFLACAKRFSPPPAME
						KCLVFEDAPNGVEAALAAGMQ
						VVMVPDGNLSRDLTTKATLVL
						NSLQDFQPELFGLPSYELEGRA
						SVFRPQPTLMVHTAGGKGKEIS
						NSSIPTCAVILAS
24026	54394	В	24164	115	594	
24027	54395	Α	24165	4	160	SIIFPAIPVSLK*IKPNCPEPPPDH
						LLQAAWGCVSLGQGLSYSALN
L	l	<u> </u>				TLLYFT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24028	54206		24166	88	2280	VPLYDRMGGETGLLIPSPFRSL
24028	54396	Α	24166	88	2280	f .
1		1				HAATDAHSCKNRHSLTGKTDS
		1				LWQQDTKLQTRPKATQGKGES
		1				GGCGPHRPLLLLRTKSTPESKS
i ,			ĺ			AHDPPVAPAQAGASGCGRGAA
		1				QKGFGAKAAVDGQGRDRTPG
					<u> </u>	VPAAGPPTSSGRGDRGPSSSRR
		1				TRSQTPESPTGRPGSRQFQAAPP
		1				PSHHSRPAHAPFPGFQMYRGPL
		1				YSSSPAPATPNTSAYESPRFCST
		}				SYISSSPRFCPTSYICSSPRFCSTS
		1				YICSSPRFCSTSY
24029	54397	A	24167	2	352	WRTLIVLGMSCLGSYPK*MHLT
	10.1337	' `	[,	_	-	HLIHCS*EARSLVTLHIIPLTICG
						EPRNIIKLVGCP*IQWKK**KKI
		1				MNSRILSPSFRSRY*TSNLLGLP
						*LRGLSPVERELKLWTNRHKLL
24020	54398	 	24169	2	252	
24030	34398	Α	24168	2	252	NKSSVFATSCSLDPISIMSLM*W
						TSVISCRSKK*SRSLRIRL*HKA
		1				KELIYPLPVRESMWGYCQLLS
						MSMPIMHAGTGEMTTG
24031	54399	Α	24169	1505	1680	RTSSLVTL*TLHLEPAGT*VLTP
						GSTTGITVSGTVLFLILLSCFLSS
						TPFRLKLACLS
24032	54400	A	24170	681	932	
24033	54401	Α	24171	1740	1950	HHHHLLQMSWSAGMQAWHW
						PSFYNVVTGKTLA/LPNLIALQH
						IPLSPAGVIAKRPAPIALPNSCA
						ALLSSQ
24034	54402	Α	24172	1	652	GKLTEARTRCCMQLTSDRHRH
		1				TALNHTTCSHHCNSGVMAGRL
		-				GPITVLSGD\GPKGMRALTMW
						HTLARPTGDLEAPRHGRPRRIT
1						ATQPDHAPPATSRNRAAHP\HN
		İ				HTP/PHPQTAH/PTATHSHLTYR
						TPRPTPATRTPL/HPHSHQQYPY
						Q/PTQRSPYLHPPPAPPHISLHPS
						TTRATVNAPDTRPTIYARESVPF
						GRLLNLGRQRGMELANILQVH
						1
24025	54402	+-	24172	267	400	GP
24035	54403	Α	24173	267	400	RFLHCGNLAGLLVGFLSFKLGE
						YEGLGPLAALTAGKGLFFLPLE
Ì						SRLRLHRRPKPRSPASFG*VSRH
ļ		 				THAIHFN
24036	54404	A	24174	422	452	
24037	54405	В	24175	131	1047	
24038	54406	Α	24176	2	199	LGLR/CARLEARARPPWATVRE
						KLWLFGHPTHMRCSWRGLFQD
I	I	1	1	1		DLGLQKPLSGGSSSAYRGCENL
		l	1			DEGECKI EGGGGGGGGTT KGCELIE

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	L .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24039	54407	A	24177	440	1746	RNLQCVFQGRIKEGIPVLALS/L RPSP/RVLHTATKQKNSCQHLE EDVD\QTKWRTDTPSMDKILME EVKLEEQLKEAVEEDKQALAD TEGSEQSSQKLVEEGNMYSIQG FCKDSLEVADVLEKATQCVPEE EIKDNNPHLKNLCETLTMSEEL SQMVDCCRLPQDVSVLGVALH WVTPQMLCVPVNEDIPEVSDT VLKAITAPPRVPIGRTSGDVTGT GVRRRVSSAMSQHRAFRDVT VTVQCLELGCFLLRVELASQFL KRPRSPESNEKVPEIEVTVEASA KMSHPTLYDATRSPWGMFQSQ SVLLSCAHHSLLRGATLDRCGS IDESEEWYGMDTFECTGVGVSI RRATEPERSSGSAAALDSHRSM HPIVNHACEGSTLRCPYENLMP DALSLSPILSSRSGDASPQAEILP VVGPFRMRRCLTVDPPGLDVT
24040	54408	Α	24178	391	656	

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SEQ ID	1 -		SEQ ID NO:	ľ	ï	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	detetion, (~possible nucleotide insertion)
24041	54409	Α	24179	1	2150	MEDSTSPKQEKENQEELGETRR
						PWEGKTAASPQYSEPESSEPLE
						AKQGPETGRQSRSSRPWSPQSR
						AKTPLGGPAGPETSSPAPVSPRE
						PSSSPSPLAPARQDLAAPPQSDR
						TTSVIPEAGTPYPDPLEQSSDKR
						ESTPHHTSQSEGNTFQQSQQPK
						PHLCGRRDVSYNNAKQKELRF
					:	DVFQEEDSNSDYDLQQPAPGGS
				1		EVAPSMLEITIQNAKAYLLKTSS
						NSGFNLYDHLSNMLTKILNERP
						ENAVDIFENISQDVKMAHFSKK
						FDALQNENELLPTYEIAEKQKA
						LFLQGHLEGVDQELEDEIAENA
		ŀ				LPNVMESAFYFEQAGVGLGTD
						ETYRIFLALKQLTDTHPIQRCRF
						WGKILGLEMNYIVAEVEFREGE
1						DEEEVEEEDVAEERDNGESEAH
		ļ.				EDEEDELPKSFYKAPQAIPKEES
						RTGANKYVYFVCNEPGRPWVK
						LPPVIPAQIVIARKIKKFFTGRLD
						APIISYPPFPGNESNYLRAQIARI
						SAGTHVSPLGFYQFGEEEGEEE
						EEAEGGRNSFEENPDFEGIQVID
						LVESLSNWVHHVQHILSQGRC
						NWFNSIQKNEEEEEEEDEEKDD
		l				SDYIEQEVGLPLLTPISEDLEIQN
		1				IPPWTTRLSSNLIPQYAIAVLQS
				<u> </u>		NLWPGAYAFSNGKKFENFYIG
						WGHKYSPDNYTPPVP\PPVYHE
						YPSVPEMAEMDDPSVEEEQAF
						RAAQEAVLLAAENEESEEDEDE
24042	54410	В	24180	1	489	<u></u>
24043	54411	A	24181	39	423	RSPTPDAASRSQCSSRRRRGVS
			_		1	KEVFVSMARTKADWPANRTG
						G*STPRKPTGPTKSPLRKSAP/A
						PTGGGERNPHRLQGLVTVAAP*
					1	NLDGYQKVHSNFLIRKLSLPSV
						WVREIASGLLKQNLRFQERQ
24044	54412	Α	24182	3	384	<u> </u>
24045	54413	A	24183	1	434	MLQNVTPARAEWQRKTALQG
		[VEDPVPARPPISSPEDLLRSGEC
						STAPTQRQVPQPVHE/SHRPTQ
						DSPGARRKPTSERGHTEQRAAP
						DREARPATRGWGLRPGGQWLG
						GSEHAAAAPGPSDHPRGRCRH
						HTILQRPRLLSGKTES
	<u> </u>	1	1	<u></u>	<u> 1</u>	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence .		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24046	54414	lΑ	24184	15	286	LPSRGAWLGTCSPPCLSLPPPP
						WAPVRPEPPR*APPPAPRRPVPS
						TTQGSLGPHCFYEL*HSPRRSA
]		İ				ASLLKPARPRAHQEERTTPDAP
						P*EL*HSP*RSAASLLKPARPRA
						HQEERTTPDAPP
24047	54415	A	24185	2	214	CLSLPPPPWDPVRPEPPQ*VPSP
		Ì				APQHPVPSTTQGLRSVGARRGT
						GRQLHLQSRCGIHWVKPAGLL
		Ì				SLGL
24048	54416	Α	24186	308	311	LPSHGAGLGTCSPPCLSLPAAP
						WAP/RVARASPTSAAPTPQHPV
						PSTTQGPLGSPSTLRKL*SFALC
						NKSCCCSLFGSTLPL*AVTLTA
						KVCSFTPKA\ARP
24049	54417	A	24187	1	669	MMNEGSGISLSGCDLGTALQLL
						SQAIVECLKLFQMLNRSTILGSG
						GQQPSSHSSTKLRPDSGAQLAS
						PSGSCTGAAGGAACQSRTVCSH
		ļ				SSALGWSMGLGAVEQGVVLV
						GEAQAAQEPMEGVGGSGMAG
:						CRSRALPHGKAAKARRAIEHST
						GGLALLGDPVHSPQPLARVLSP
						SLPRASRAGW/PAPSAGPPSPHP
						PGTPAGPQAPHAAPVPARASPS
						TPPCKL
24050	54418	Α	24188	157	395	RCSSCQREMTPYGK*NVQKRL
						KTTKNGSM*LKSHILSTPEQ\PA
						SPTSS/PPLLQPAAPAPPLMPRPS
		1				PPPGISVA*YPGM
24051	54419	Α	24189	2	287	CLSLPPPPWAPVRPKPPR*ATPP
		1				APRRPVPSTTQGLRSAGARRGT
						GRQLHLQPLCGIHWVKPAGLLS
						LVGTWRTFMSSSGIVNTPISTLC
						LAQGL
24052	54420	Α	24190	500	724	
24053	54421	В	24191	1	2154	
24054	54422	A	24192	688	1215	SRLAPALLSPWVVDGTGRCGA
						VGGAHWGGLGRTGAHGGGGR
		1				LRHGGLRVPSPAPREGS*GPPR
						YRAQRWWAGT/DWRTQYTLRS
						CWPGTLPSK*REMTPYGK*NV
						QKRLKTTKNGSM*LKSHILSTP
			<u> </u>			EQ\PASPTSS/PPLLQPAAPAPPL
						MPRPSPPPGISVA*YPGMPWTC
						PFYWQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24077	15.400	 	0.4100	la on	[1121	W. O. I. P. P. T. WOOD, P. A. WOWEN
24055	54423	Α	24193	308	1131	VLQLLRRRTWSCSLRAWSWDL
	1	i				IVPPGGLVVSLASGVKLQTFAV
						SVTALKEARLDLFVPPGGLVVS
						LASGVKLQTFAVSVTAHKSSV
						DPKNSGAQLASPSGSRTETAGG
ŀ						AACQSRAVRSHSSALGWSMGL
ļ						GAIEQGAALVRGGSGRTG\PME
		İ			Ì	WVGGSGMAGCRSRALPRGKA
						AKAQREIEHSAGGPALLGDSVH
1						PPQPLARVLSPPLPGASRACWL
		1				LRVRGPPSPRPPGTPAGPQAPH
		1				AAPVPARASPSTPPLVHLTNDV
						LRVKVKWAFFFTLHK
24056	54424	Α	24194	192	818	SWCLSRAVVGDSMTSEVIED\E
		1				KQF\YSKA\KTYWKQIPPTVDR
						HAWGEGPNKTGTSCALDCGAG
						IGRITKRLLLPLFREVDMVDITE
		1				DFLVQAKTYLGEEGKRVRNYF
ļ		İ				CCGLQDFTP\EPDSYDVIW\IQW
						VIGHLT\DQHLA/ESFLRPCKGQ
						ASAPNGIIVIK\DNIA\QEGVILD
						D\VDSSVCRDLDVVRRII\CSAG
1		1				LSLLAEERQENLPD
24057	54425	Α	24195	1	233	FFFFETESRCVARLECSGAISAH
		1				CNLCLPGASNSPASASQ/CSWD
		į				YRCVPSRLANFLYFSTDGISPC
		ļ				WPRMVSNLLTS
24058	54426	Α	24196	240	476	LLPQYRRVLLIYSGIQLLPGLGL
						RRRMCPGIYSFLLDFLVYLRRG
						VYSIL*W*FVFVVLVPWFSAPS
ŀ		Į.				GHLRSSLCCLF
24059	54427	A	24197	288	677	NLPEDVVPDDKPHQPLLRAV\C
					}	LFVISKDLFLESPFESPIWRMYP
						NSTVHIQRLELALLC/CHCPRAL
		ŀ				PASAHSCHSLERGLPICKE**CM
						PCSEYTIASQRPAGGAPAILPEC
						DSRRRTTQATSEPGSHI
24060	54428	Α	24198	99	399	LLCHGFEHPPLAQRSLLLPIV*S
						LLLSTRQSHSPSSFVPLLARSCG
						SWVSPAEAAEPQILQKGKCSCL
						IVPLEALCQRGTRLSLWLPLTFF
					1	PSFQLWPV
24061	54429	A	24199	1	586	LSFFKVFSFFAIGSNILFSSEKFII
		_				TDHLKPSSPTCQSHSLSSFVPLL
					1	ARVAFRWGFGVDVLSFVSFPSN
						SQDPQLQVCGVCWRSTPDRLP
						GYHQRGSGGCRTADIIEPQMLR
						LIVRLEVLSQRSTRPQRCQ\SA\L
						LGGASQLGCC/GVRDLLEEAVC
					1	PFSDL/KLHAGKTTTLFKAVRQ/
						SHLSLQRLLLSFVCLCPT
			<u> </u>	<u> </u>	<u> </u>	DITTOT CICETOL ACTOL I

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24062	154420	1	124200	1	822	
24062 24063	54430 54431	A	24200 24201	1428	1801	RSAGIDPE*FFSG/PRRIHTASCL
24003	34431	^	24201	1420	1601	PSQRSSNRACSSSVTRIVSILSRF
						IGIITPMNRNPRFWRNSTSWTR
						MNRQTSVNRFTTTLMSFTAAA
						SRVSVMTVKTSDTCSSRRRSQL
						VKYHLSISTQVL
24064	54432	В	24202	1	2040	
24065	54433	Α	24203	176	564	AWEPKWYSTTATRTHFWEMRS
						RSIMSTTKAFSTSNSASDSLEEP
						SSRKTRSSSLLQ*KSCNGGSEM
			Ì			SSTYRVVGLTGASLSASALNSLI
						DSRSNR*V/FSRNTLANITRRNQ
			_			LKMLPAPDPDGFSSGSPFQ
24066	54434	Α	24204	1	774	
24067	54435	Α	24205	1	1807	MPLRAFNPVSSFRWARGMTIV
						AALMTVFFIMQLVGQVPAALW
						VIFGEDRFRWSATMIGLSLAVF
						GILHALAQAFVTGPATKRFGEK
						QAIIAGMAADALGYVLLAFAT
						RGWMAFPIMILLASGGIGMPAL QAMLSRQPNFDHWTADRHGD
						LCRLGEHMERVGMDCRRRPIPC
						LPPRVASRCMEPGHLDLNGSRR
:						HLANGFTTPRIGANQFLRRTVN
į						AQTNPWQNISIASAISSSRTRRIS
:						GSVGSWPRSGNAEVSALHHYV
						PDLHRRMLLATLWNTYICINEA
				•		LALTLSDFSLVPPHPYRQLFTLT
						TFHNRACSSSVTRIVSILSRFIGII
						TPMNRNPPYTEASVTKQEKTAL
						NMARFIRSQTLTLLEKLNELDA
						DEQADICESLHDHADELYRSCL
						ARFGDDGNSYDHDYAKLACLQ
						VEGRGGGTFAHELLAVEYAGW
						ISPAFRLKVMHWLMNRMLDAR
						NRDRYKQWKQDFFHDSYNRQS
					:	AGHILSQCANLAATTSEYFIHKP
						HRLIAAETGYSQSTVVRAFREA
			1			VNKGILSVEIVIGDHRERRADL
						YRFTPSFLVFAQQAKNALIESKS
						KI\SSAATKVRLLAPISPTSPMG KIGLATSGS
24068	54436	A	24206	291	851	RIOLATOUS
24069	54437	A	24207	2	694	
24070	54438	A	24208	2	331	HLSTIEPEDYNLLFRRVSLKLSV
[Γ`		<u> </u>		WLPTWQSKSNGWSKKCLWMH
			1			LKSNPLNALLPPAGF*SATMPK
						CSAWAKTISSPGCVITEF*SQPV
						NAGTSPNKNTYLVGISPLKKP
24071	54439	Α	24209	1	4455	
		*		• • • • • • • • • • • • • • • • • • • •	•	'

ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
of peptide	hod	-	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
54440	Α	24210	698	1088	LTSKFIAAVCMRRISLCGKGSA
					QNTDSGIAATPCSCVSD*QNSTS
	İ				SRIRM*PWMCPCPMKRKSAT*N
					ITLRC*RAAPPISSRVRK*/MSM
	1				Y/NHLQSAECGGHRPFPVPQEE
1	İ				SILLLHLIANRSSACHSIR
54441	Α	24211	1	1902	
54442	Α	24212	1	2898	
54443	Α	24213	1	2604	
54444	В	24214	1	1254	
54445	Α	24215	1	2946	
54446	Α	24216	698	1285	RFWRNSTSWTRMNRQTSVNRF
		l i			TTTLMSFTAAASRVSVMTETA
	İ				MTMITPSLHACRSKAVAAALLP
					MNCSLWSTQAG/CSP/DVSAEG
					KPDIYRLSNRKITTCYSAES/LL
					KLSVWLPTWQSKSNGWSKKCL
					WMHLKSNSLNALLPPAGF*SAT
					MPKCSAWAKTISSPGCVITEF*S
					QPVNAGTSPNKNTYLVGISPLK KP
54447	Α	24217	1	1986	
54448	Α	24218	1	2262	
54449	В	24219	1	1014	
54450	A	24220	1	1155	
54451	Α	24221	5	386	
54452	A	24222	1922	2146	SSSAGISCWSSSPTPGSIPSRVGS AAA*GWTTSRSSTGSANPSASR KPAAAIRASMPPNCRSGEARWP LWSGSL
54453	A	24223	1091	1435	
54454	Α	24224	1	2919	
54455	Α	24225	1324	1502	
54456	Α	24226	1	2901	
54457	Α	24227	5	253	
54458	Α	24228	1227	3031	
	54440 54441 54442 54443 54444 54445 54446 54446 54450 54450 54451 54452 54453 54454 54456 54457	54441 A 54442 A 54443 A 54444 B 54445 A 54446 A 54447 A 54448 A 54449 B 54450 A 54451 A 54452 A	of peptide sequence hod in USSN 09/540,217 54440 A 24210 54441 A 24211 54442 A 24212 54443 A 24213 54444 B 24214 54445 A 24215 54446 A 24216 54448 A 24218 54449 B 24219 54450 A 24220 54451 A 24221 54452 A 24222 54453 A 24223 54454 A 24225 54456 A 24226 54457 A 24227	sequence 09/540,217 codon for peptide sequence 54440 A 24210 698 54441 A 24211 1 54442 A 24212 1 54443 A 24213 1 54444 B 24214 1 54445 A 24215 1 54446 A 24216 698 54447 A 24217 1 54448 A 24218 1 54449 B 24219 1 54450 A 24220 1 54451 A 24221 5 54452 A 24222 1922 54453 A 24223 1091 54454 A 24224 1 54455 A 24225 1324 54456 A 24227 5	of peptide sequence hod 99540,217 in USSN o99540,217 location of first codon for peptide sequence codon for last amino acid of peptide sequence 54440 A 24210 698 1088 54441 A 24211 1 1902 54442 A 24212 1 2898 54443 A 24213 1 2604 54445 A 24215 1 2946 54446 A 24216 698 1285 54447 A 24217 1 1986 54448 A 24218 1 2262 54449 B 24219 1 1014 54450 A 24220 1 1155 54451 A 24221 5 386 54452 A 24222 1922 2146 54453 A 24223 1091 1435 54454 A 24224 1 2919 54455

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u> </u>	<u> </u>				
24091	54459	Α	24229	1	1940	MNRQTSVNRFTTTLMSFTAAA
	ľ	-				SRVSVMTVKTSDTCSSRRRSQL
1						VCKRMPGADKPVEGRGGGTFA
						HELLAVEYAGWISPAFRLKVN
•						QTFIDYRTGRLQPAIPQSLPEAL
						RLAADLAEQKQRLEQKMLMD
		l				APKVEFAERVATASGVLIGNYA
		l				KVLGLGQNYLFTWLRDNGILIA
						TGERRNVPKQEYISRGYFTLKE
		ļ				TVIDTSNGSRISFTTRITGKDVV
		į				RTTSIVMLAKVTFLSCITMSDFT
						FSGYELACFVTHSGLSRSAGHIL
		ł				SQCANLAATTSEYFIHKPHRLIA
	<u> </u>	ŀ				AETGYSQSTVVRAFREAVNKGI
						LSVEIVIGDHRERRANLYRFTPS
						F\LAFAQQAKNALIESKLKISSA
						ATKVKAVLAKTLALFNFLSTPP
						CQNDTPSPCQDDVAIKNKKSQ
İ						VKKTKRSVSGGAGTTSLKKLTS
						WIAKAKAKADNLRLSKKRTQK
						- 1
						HEFKQKVEAAARKYAYLKNKR
						SPDIGGISNFDNLPHCMTVNEA
	:	1				LNAVLAKNKDNEQWGAVAGA
		[YIADITDGEDRARHFGLMSACF
						GVGMVAGPVAGGLLGAISLHA
	ŀ	Ì				PFLAAAVLNGLNLLLGCFLMQ
						ESHKGERRGELLKIVVLPGDHV
						GQEITAEAIKVLKAISDVRSNV
						KFDFENHLIGGAAIDLQGGGGK
						PRCVSKSLMLHCTR
24092	54460	Α	24230	1	1230	
24093	54461	Α	24231	1	799	MKPRTYTVSVTAVKDGVSGVC
		1				SFRCSDVSRVSSFQWVRGLADF
		ŀ				RSEAADLHNSGAQLASPIGSRT
			ļ			GATSGAPCQSRTHAPALLSLW
		ŀ				AVDGTGRHGAGGSTRRQGSGG
ĺ						TQEPMVAGKAQAWRAAGFSG
Ì						SGDGVGVSILVPWGGTCIGSFR
						QANSWISGGAFSDAGCDSDRL
						GKDWNSLEGSEEERKMWESLE
						LPRDLKSSEDRKMWESLELPRD
						LLNGFDQNAD\TICTMKSRLRW
]				SQMEMKNLLGTGVNVTLVML*
		İ				QRDWRHLLLV
24094	54462	A	24232	416	775	
24095	54463	A	24233	548	882	PGGHQYPPCHLEEKETDLPLLY
		l' -	255			TAVEEKGLSYSIIPGSLVEKPLT
						STNSSGTPIQSNITSL*IPLAGIMS
						MAASQN*AI*WPLAPTMQSFTC
ĺ						TT*RLS*SHRRSWWHRGCT
24096	54464	Ā	24234	144	570	11 ICO SHRISW WIROUT
24070	124404	<u>Γ</u> .	44434	1177	1310	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24097 54465	A	24235	254	1117	LSRMKLRTLVVSVTVLKDVVS GVCSFRYSDVSGVSSFQWVRG LADFRSEAADLRGLKLQTFVVS VTPHKGSVDPKSDQQRDLLQR VKEQNFRNVKGDPAALAHEDD	
						VNGTESGAPRKHGDRELDDVV ITNPHIEAILENKDWIEDASGLM SHCIAILKICHTLTEKLVAMTM GSGAKMKTSAIVSDMIVVAKW NSPRVDDVVKSMYPPLDPKLL DTGTTTLLLSV\SHLVLVTRNA
						CHL\TG\GLDWI\DQSLS\AAEEH LEVLREAAL\ASEPDKGLPGPEG FLQEQSAI
24098	54466	A	24236	472	844	RTDGVLVWMSFLFVSFPSNSQD PQLQVCWSLLEVHSRPCLPGYQ QWRLLCPHEITGESMSKLHIIQK YTSSCCLFKSS\SERGSSTPCPSW LGSGEGLSTLHATDFSFSAPMA QRGKSSLVSLE
24099	54467	A	24237	206	456	
24100	54468	A	24238	50	392	LGLLPFFHRCPTQRGGI*RGSLA TAALLSCGGLCPVGS\SRHLCL\ PVRGKRPTQASVMGDALPPTK LECPRSASGCCAGSDNFKPVDL SLLSSMGLGSTELDHLAPWLQP PFQG
24101	54469	A	24239	1	204	SFSVQFCSLVGDPLG*SSCDPLE EKRCSGFWNIQPFCAGFSPSLW DYLSLVFDVDDLQMRSLSGCPF
24102	54470	A	24240	1	375	MCNAQVLEYMGKSSSLLTSDL QLVRDALRSLRNSFSGHDPQHH TIDSLEQGISSLMERLHVMETQ KKQERKPPLVIPRQT/WVWSGP PANSNRPAAEGPDC*KEN*RDA LRSLRNSFSGHDPQHHTIDSLEQ GISSLMERLHVMETQKKQERKP PLVIPRQTGSGVDLQQTPTDLQ LRDLTVRRKTNEQKGIASASTK RTSTPKPHL
24103	54471	A	24241	1	674	MVVYACSPSYLEAEVGGSLSLE VEATNPKNSSKKFLDLISEFCK VSGYKISVHKSVALLYTKNDQ AATQIKNLIPFTTAAKQFKYFGI YLTKEEERLKPSRKKITKKHTK KRTASLILHAMICSQIPKQQNE KYQVPQFDQSTIKNIESAKGLD VWDSWPLQNADGTPPLVIPRQ T/WVWSGPPANSNRPAAEGPDC QKEN*QTERNSINIKKKDIHTKT PLCI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24104	54472	Α	24242	1676	1768	
24105	54473	Α	24243	270	471	PSREYQDANPWLKWIKYSVRT
						LNKSDCYASGYK*ACLLCSVYS
						HGKAAGECTLSAGSKNGLAEVI KK
24106	54474	Α	24244	23	200	AQMKAHGRSRSS*RC\DEEHWR
						ACRAQEPPCSQSQRERSKRTCR QGTVASKVKCHSEAR
24107	54475	A	24245	3	270	ASTSHVPQPICGLDDLAIRHSAV
						RLNRERVHSTAVLP*EYTKQRR
						QGHL*PDPYKKEGAATNEDAL
						LLRWERGRGARLHIPGCKAPQL
24108	54476	A	24246	135	275	
24109	54477	Α	24247	365	583	
24110	54478	Α	24248	2906	3114	
24111	54479	Α	24249	108	405	NLHTPYRPQSLGQGEPLDDYV
						NTQGAS/LFSVTKKELGAGSIEE
						CAAKCEEDKEFTCRYFHCRCTF
						LSIPQVTSSFLCFPLPLPLLKSSK
						LLANRDKYRM
24112	54480	Α	24250	.1	386	
24113	54481	Α	24251	1	808	MSYSQCVSDPKAHGILPGYHC
						WLFRDQGLLKSQHVINPAKAST
1						GFFLQDTLMVLEIKTTIRNYYK
						HLYAHELERLEEMDKFLDTYT
						LRRLSQEVIDSPNRPIMSSKIESV
						KGYTHQQFCHKSTLNKGDRVI
1						DNLTCPPYCCVWFPLAGTGPHI
	1					LYLSRLVSNLELFKRGKGRGEQ
	}					RKEEVTCGMLRKSLVTSDLPEP
						GGVTSEGCETAKVTACSSLWE
						LCHREVLTPYCPKDTCRKWLE
						TPVRRSCPVRRSRISTCLRKKFG
						HILAEQLSCA*RCL**FL/YGCFY
	1					FQDHECVLKEESSTSFGRINHV
					_	LTLKEPLVPE
24114	54482	A	24252	833	1245	ANMFPAFAAVAAGAATEAAA
						AAAAAPAPLAPARRAAAAARG
						PPLPRLP/WSCAEVMYHPQPYG
						ASQYLPNPMAATTCPTAYY/RA
						GAPTWPAGESPDGQPAGGGGE
						GHREYRSGARRRPAGTRGAED
				}	1	ALGSAGASLSRKP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	<u> </u>	<u> </u>				
24115	54483	Α	24253	145	766	TGCIREKFHLPEGSLLIVSSRKH
		ì				LQRLAFQAKVSSMMLPLQGAQ
						MLQMLEKSLR\KSLPRNPLKVY
		1				GT\VFHINHG\NPFNLKAVV\DK
						W\PDFN\TVVV\CPQ\EQDMDRW
		1		İ		TLDHYTIT\YQIYSKDPQNCQEF
1		1				LGSPELINWKQHLQIQSSQPSLN
						EAIQNLAAIKSFKVKQ\TQRILY
•						MGRLETAK\ELTPFLAGNQRILS
		<u> </u>				P\SGWAKPKAM
24116	54484	A	24254	215	550	VGFRSCLVMSGRGKGGKGLGK
						GGAKRHRKVLRDNIQGITKPAI
		1				RRLARRGGVKRISGLIYEETRG
		1				VLKVFLENVIRDAVT\YTEHAK
						RKTVTAMDVVYAL\KRQGRTL
24117	54485	A	24255	3	292	TLLNLAAAHCSGPCLLWFQLSF
						RSPSTSAVGRCRRPSADFHPSRS
						SRVSAVLLIQ/QRCPLPILIQLKA
						CHCSCTAKWLGSSPKPETFFEII
						HPSPS
24118	54486	Α	24256	1	789	
24119	54487	Α	24257	3	256	TLLNLATALSSGPCLLRLELSFH
				1		SPSTTAVCCRHRPAADFPPFGR
]						VSSLLLIQ/QRCPLPLLIGLKACQ
						CSCAAKCCVRPNRAEQ
24120	54488	Α	24258	3	212	
24121	54489	Α	24259	2	491	
24122	54490	Α	24260	20	272	
24123	54491	Α	24261	160	876	
24124	54492	Α	24262	623	796	GANSWNLTGTLAVKAGTVENP
						I*VFATHVESAGPR*LDPRTGSL
<u> </u>	ļ	<u> </u>		1		AVRTKGRHWDQGA
24125	54493	Α	24263	242	659	TSNSPPQTRPHVCHGPRQYLLQ
						LSSRDLEFLVQIAVRSCLCPGLR
						IRSPVFAVASTSTFGTLRGKVV
ļ						KDFGFVFPPPWGQPCLCPVEER
						ALSCQGPVVDSRCPRCGHEGM
						AYHTQQMRSADEGQTVFYT\C
		1		1		TNCKFQEK
24126	54494	Α	24264	66	238	
24127	54495	A	24265	2	278	TENTED AT COOR OF COMMON O
24128	54496	Α	24266	168	565	TENFRALSTGEKGFGYKSSCFH
		1				RIIPGFMCQGGDFIRHNGTGGK
						SIYGEKFDDENFILKHTGPGML
						SMANAGPNTNGSQFLICTAKTE
						WLDGKLVVFGKVNGMNIVEA
	<u> </u>	1		<u> </u>		MECFVSRNGKTGQKITIA\DCG
24129	54497	B	24267	20	88	
24130	54498	С	24268	183	254	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	1	Į.	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24131	54499	A	24269	412	590	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE KRDLPLRVKDQGRRPCVV
24132	54500	A	24270	550	852	
24133	54501	A	24271	207	381	
24134	54502	A	24272	412	589	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE KRDLPLRVKDQGRRPCVV
24135	54503	Α	24273	191	235	
24136	54504	Α	24274	95	375	AEKIWEGVSQRALGQSSRGSGS GCQMNPLSGRPQVVCFFTAHM LTKKREEGASMLNMPCFQ*KE EMIQANLGPIADSSHCLGIEGPP WAPLTP
24137	54505	A	24275	144	293	WQVPLFWQPFPEPLELWPKAL* LTPSQIFSD*RLKTDAARSPRKP PAPR
24138	54506	c	24276	1	909	
24139	54507	A	24277		1744	MGCLGTQRAPRAFLLLPLPLHF AQLSNLTQLQAAASLSMVTHR SSGGGGERGATAAGTEQGGAC LAASLLRGLWDAGEVTPIRVRV RPYDPFTLAPLHEHSEAQCVAS PSLTVTKEKSLLPKSLLAGSREV ARSRALCHGQARPPPRPQVTQQ RTQLQGQRILQWRPATEGPHSP RRGSDSAKGTHPPTTGTCPPRR RQGAGYSRYNSGESRRLGRQ QDLVARSDEQRDPAPATRRED KMIEELQPFGEHRPRRSPSQGC DTLFQALWFLASPSFQRGGQLQ VLALCKAAAGPGAPQAASTAG TREHGYAWKLGDTRNHRALK RKSQTWLRRFPAGCPDVCSAL ADLRAFMGLRGEEVCADWSM GGHGSGECRRGTGGQRAVVPK SPVQEKGKEPLSLLQYFMVDEK ERFLRITSEHEDNDNSNNSSNHS SINETKETCFIHEPKTPAPVTDW EGSLPLVFNHCRDASLIIHSCFK GVRPHRDACLG/RFTLSGK/FPL SWGRGKYPSTPSPSPLAASPTFL GQGQELVTSARNLTTRPRNAC GPGFLLSHVPSVRDPTGNRTVQ
24140	54508	A	24278	2	120	LTWQPLPEPLELWPKAL QSSAAKPRTSGKGSVREPWAR VPGALGVAAR*AEQSDFQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24141	54509	А	24279	1135	1483	ILTMTGSLSCTPSNAVIEKV*VC CSSLSTG*SKV/QVASNTSLCDG ED*GVVIQKKVKLWVFCIFYVL VQSYYS*R*SQIETVRNGCSHW GRFQDIALITGHCKIWTLIVFIQ YFNV
24142	54510	Α	24280		860	ATTGGTNLRDDIMRLDDTVHV VIATPGRILDLIKKGVAKVDHV QMIVLDEADKLLSQDFVQIMED IILTLPKNRQILLYSATFPLSVQK FMNSHLQKPYEINLMEELTLKG VTQYAYVTERQKVHCLNTLF SRLQINQSIIFCNSSQRVELLAK KISQLGYSCFYIHAKMRQEHRN RVFHDFRNGLCRNLVCTDLFTR GIDIQAVNVVINFDFPKLAETYL HRIGRSGRFGHLGLAINLITYDD RFNLKSIEAQL*TEIKPIPSNIDK SLYVAEYHSEPVEDEKP
24143	54511	A	24281	848	1218	PMQRKSMLKISVS/PVSVQKFM NSHLQKPYEINLMEELTLKGVT QYYACITELQKLAETYLHRIGR SGRFGHPGLAINLITYDDHFNL KSIEEQLGTEIKPIPSNIDKSLCG AEYHSEPVEDEKP
24144	54512	Α	24282	241	477	
24145	54513	A	24283	38	208	MTAGAPPPPPPFET\PEGV*GNP TCSGG*EDASGSTRSQRHPVGL PVSCALRACHLK
24146	54514	A	24284	794	1455	AQQGGTSKKRRFPQDGQQRAG LACSRPPGPGGCGFQAPAGERY PNIKPHPTSVAPLVLCICYLM/P MLPWAWSYWRGARRPSRHPG AFPCSVAHQCPSAARSPTLRQE DPEKGSQASSFPTGPSPQSMVD RCLKGPQLDVSRKQLSLPQTGL RPVLVNQLSKKPTGYPQTQIQW VLMRDTGKDTDTEEKEGHAK MEAEIGVMYPQEHLEPPESGRA KGEIPL
24147	54515	A	24285	90	354	QSPGVHQLSRFVS*SYSERP*SL LVSTRYHTRALH**HYADSIQ* ARSSKHTGLIGCTVP*RGFPRTH SFYSRKREPRRQTSSFRGIL
24148	54516	A	24286	236	520	TACFLTHVAIKWVMHSSIPSSN GGGIYMIGLKAQVSYMRKWLK SPWSPLLLSPSLHRWPHGKFPM IS*QRKRRLGPGSQMVLHDMQ APPKSGQL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24149	54517	Α	24287	16	278	FAQAMKPHLVQQLQLESPLG*
						AYDNPLSFSKIHLSDLHKPVL*L
						GDSNDVLGTLESMSAQSQCPV
		l				VPEMSDHLPFPMHSYPGKRPEV
24150	54518	Α	24288	74	168	LGKCLFLHSCP*APPEAICLQLP
		_				LLSSPNGP
24151	54519	С	24289	92	564	
24152	54520	Α	24290	1	876	
24153	54521	Α	24291	71	282	PLVTQVVFRLFPTNSSLDLLQ/S
						PSP/GSGPHSNWQPFGSLDKWA
						GVKHPNEECAASKIQTGPPCVV
	1					PLPWL
24154	54522	Α	24292	81	1054	PRLFSDCFPPTAAWTCCRAFSC
						SGPHSNWQPFGSLDKWAGVKH
						PNEECAASKIQTGPPCVVPLPW
						L*EEPNAAIYPSPVSPVCFLLLA
	1					HWIQSA*CHQCNGPVWYLVEV
						KSNQVLSK*HYDTKPES*YTPY
						GQEWRKGHLPSQWPHTRYQE
	1					MY*FAQAMKPHLV*QLQLQSP
						LG*AYNNPLSFSKIHLPSPHGSH
						SEPCL*GPTRTLV*L*L*KQTGM
		l		1		LAVAPEENQHYLAWQLPQGRP
						SLLPQAVQGLSPQTKVERLMA
						ASVGEGMLPILGMEKLF/PSGK
						KD*SGFTSSVSPASSGSSHTSPF
						QVAGSHSFPINAFTLTVDTW
24155	54523	Α	24293	5	88	
24156	54524	Α	24294	779	1041	PPLSSPSGP/DEQSD/LWWQQSIP
						LHQETVKVLH*MGPVPCATQL
						DKIHQQGFSDPYRGAVLLASG
		1				WCPKRPEIPEEGATTHLCCFPAS
24157	54525	Α	24295	248	3714	
24158	54526	Α	24296	4177	7804	TFQESRFEQGFQKAIFILELVDA
		1				EDPPDEEEEEASSIFSSSFHFLFP
		1				SSSSCFSFSSSSSSLLLRSPGDKD
		1				MPTAGMPSLLQSSSESPQSYPE
						GEDSQSPLQIPQSSPESDDTLYP
						LQSPQSRSEGEDSSDPLQRPPEG
]						KDSQSPLQIPQSSPEGDDTQSPL
						QNSQSSPEGKDSLSPLEISQSPPE
						GEDVQSPLQNPASSFFSSALLSI
						FQSSPESIQSPFEGFPQSVLQIPV
						SAASSSTLVSIFQSSPESTQSP
24159	54527	A	24297	145	219	
	1		<u> </u>	1	J	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	j	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24160	54528	A	24298	191	1075	QPKVSRHCQTSPGSYLYSLQSG
2.700	3.320		2,250	.,,,		RWTTGKPLQQTKSVYLRSFHH
						HHFQNHRHYLHHHHLYHHHH
						ҮНРНРЅНННЦНСҮ\SQHНННН
						HHHHHHHEHHLCRHOKSTOVE
						YSVFAERENDLGLNGTGDEAL
	ļ					HDNGTTERVSGPPAGYLYRAD
						YAIGLQPFMPRLGRRTPGAGGP
						HSRAPELAYLQKLITSPATLPFV
						YLETGGWSPRDAHRVDNFDM
				·		GPRENMGSVWEKSFLFTRNPSK
						SSYIADKNLIYLYASECIGRQSQ
						KARASKSGVQAEHPYITGAIKD
						DTRAIVKTRDLANRLK
24161	54529	A	24299	155	381	LICTSFKILSRLFCPSSL*SSSSQG
21101	3.323	`	2 1299			ACISKLLSVLTYAHLYFALEKE
						VLTFLSRAIVSSSTLLTHGVAHS
					}	LTLFH
24162	54530	A	24300	1052	1338	TGWDGPPANCSSPMEGDSNVH
		-				CSCVCCCSCFGLLL/CAIHCCCA
						PPGCSACGAATLLYSAPLGSSA
						YGIALLSPFHHVRMMLSTKQEV
						GLHQTLNLP
24163	54531	Α	24301	1	723	MLNVQALVVMVMCFMWASL
						WAPGGHPSAGNISSWVSGLVSF
						PEWSSWELVVSVVGWAGKLW
						LHESLLTLNLSCGLLVVLVALV
		l				FDEWHPWSGHVAEACSTGYWS
						TGGIWFHEIAAP*DCSSSPAME
		ł				QSWMEND\DELIEVGFRRLVIT
		l				NFSELKEDVRTHLKEAKNLEKR
		ł				LDEWLTRINSIEKTLNDLMELK
	1					TMARELRDACTSFSSQFDQVEE
		l				RVSVIEDQMNEMKREEKFREK
						RVKRNEQSLQEI
24164	54532	С	24302	295	430	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO:	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24165	54533	Α	24303	811	1947	QTERNSININRNGHQHQNPICRS
ĺ						PSSKTKGSIKPQRSGRNQSRKA
						ENSKNQSTSSPPKDCSSSLAMK
		,				QSWMENDFDELTGAGFRRLVI
i	<i>'</i>					TDFSELKEDVQTHHKEAKNLE
						KKIRQM\VTRINSVEKSLKDLM
				į		ELKTMARELCDACTSFSSRFDQ
	1					VEERVSVIEDQMNEMKRKEKF
						REKRVKRNEQSLQVIWDYVKR
						P\NLRLIGIPESDEENGTKLENTL
						QDIIQENFPNIARQAN\VQIQEIQ
						RTPQRYSSRRATPRHIIVRFTKV
		İ				EMKEK\MLRAARQKGRVTHKG
						KPIRLTADLSAETVQARREWGP
		1				IFNILKGKNFQPRISYPAKLSFIS
						EGEIKYFTDKQMLKDFVTTRPA LQELLKEALNMERNNWYQPLQ
24166	54534	A	24304	1	2394	EQELEKEALIMEKINW TQFEQ
24167	54535	A	24305	1902	2413	
24168	54536	Α	24306	2	610	WAPPAIHIRCMPEAYNFPRKPL
						SLIPSAIQSNELIIMNGNRALNIQ
						LLKRYQCTLDIRYKQTSMLNC
		İ				YEGTVITPSHIQYPCLSDSYGLY
		:				DIGIRNTDFLKGELSMNIRPLHD
		İ				RVIVKRKEVETKSAGGIVLTGS
						AAAKSTRGEVLAVGNGRILEN
					,	GEVKPLDVKVGDIVIFNDGY\G
	<u> </u>	1				VKSEKIDYEEVLIMSESDILAIV
24169	54537	A	24307	1	801	TA INCORPORATION A COMPANY
24170	54538	A	24308	1268	1579	TAIWSPSKSALYAAQTSGCSWI
					:	ALPSISTGSNAWIPRRCRVGARF
						SSTGCSRITSSRISQTTASSRSTIF FAALMVVARPRSSSLP*MNGLN
į						SSSAIFFGRPH
24171	54539	A	24309	583	796	IHICDIRFSATPAGGEFWQYIAS
241/1	37339	^	24309	1303	1790	ROSIFHAIVETGHALIQPSSVPG
						LTPAMKAVATTSH*TVESCFRS
						ASR
24172	54540	Α	24310	1	313	MPNVDVVGEMVNTMSASRSY
						QANVEVLNTEKAMSIAVTTTDP
						TNTGVSTTSSSSLTG\SNAADLQ
						SSFLTLLVAQLKNQDPTNPMEN
						NELTSQLAQISTTLRAY
24173	54541	Α	24311	54	323	SASAQVSALSIHSAAFQNEYLSP
						CPN*RLV\SAFTNSSRQSG*PEK
						SVSQTPVITASAFT*YADIAASV
						RNKILRPGTNVFGDTIGFRVQ
24174	54542	Α	24312	816	866	
24175	54543	В	24313	1	985	
24176	54544	A	24314	1828	2505	
24177	54545	Α	24315]1	1938	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24178	54546	Α	24316	4201	4668	
24179	54547	Α	24317	269	503	
24180	54548	В	24318	1	1702	
24181	54549	Α	24319	829	1584	
24182	54550	Α	24320	1238	1926	IQAAYNRIAPSSSPKFPRACGKP
						CLGERGRIRGPGPSFGLINRPST
						PDPTITGVSTT/SSSSLTGSNAAD
				İ		VQS/SFLTLLVAQLKNQDPTNP/
l						MENNELTSQLAQISTVS/GIEKL
						NTTLGSISGQI/DNSQSLQASNLI
						GHGVM/IPGTTVLAGTGSEEGA
						V/TTTPFGVE\WDGTL/TDGST
						APNGSYNVAIS/ASNGGTQLVA
						QPLQFALAAERFCALSPGAPYR YDTWVRQWVASP
24183	54551	A	24321	487	1509	FYLOGLROOPQVROPLNIGVTA
		1				QGVNAAACHPHIAEQQLDHRH
		1	ļ	3		RTNVLRTNGVLRPAEGIQERGS
		1				FIVSTGFCDVFADFQERVFWRT
						TDVFNHIWRVAGNVLFQQVPH
		1				AARMLQRGIAFGEAVFVQFISP
		1				GGFIVLAFFRVVAGEQAIFKAVI
						LTHDQAGGGIGFGVFAVEFFIV
						QQVQQYA*QESNVSTGTNRGIV
						VSYRRTTRKARIDHNQLRAVV
						NFGFHRPAEPDRVSAHHHDEV
		1				GVLDVDPVVGHRKTFFDIWHD
						DQFIDQRVRRFSGDNRRLQDRE
		1				QVGLQSANNSPIDKSLFRGINQF
		1				QLYAAFTAQDVNIDIFKTRQQF
				_		FAVIGQTTGVQYGK
24184	54552	Α	24322	377	749	
24185	54553	Α	24323	140	298	
24186	54554	Α	24324	342	1094	MGAPWLHRPRRALQRWRLGS
		İ				GFSCGSGLGLQENNGGRPVMV
						LALSTLPRRPTRCGCEFYCLIKP
						GFRLGMKSVKMEIRGLWAGGE
		ł				DKIAVPAEPRSAGPERTLSDLH
		ł				NESIF/ITGGGSGLGLALVERF/IE
]				1		EGAQVATLELSAAKV/ASLR/Q
				1		RFGEHILAVEGN/VTCYADYQR
						AVDQILT/RSGK/LVNTPAETLE
						TGFHELFNVNVLGYLLGAKAC APALIASEGSMIFTLSNAAWYP\
						GGGGPLYTASKHAATGR
24187	54555	Α	24325	1	1785	· · · · · · · · · · · · · · · · · · ·
24188	54556	Α	24326	2	1760	
24189	54557	Α	24327	2437	2679	TFQYIGQPHGDEFPHVAPVKHG
						DKLGEGKNHHDKSRHLFHRFN
						H*MR*IRLTSNCAGSDAISQRD
		1		_		NKEKKNDQWHDTFNKG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24190	54558	Α	24328	3	415	
24191	54559	A	24329	1	1932	
24192	54560	Α	24330	1932	2117	RAISLGRTAAGYLNSDVAQRA
						ASGADPAGDFVVWH**IRKNLS
						GGARYAIPHLYQHLAWHP
24193	54561	Α	24331	1	1017	MRYREPPARGVEPNELHACRSE
1						DKLSEDAFDDQCTGANPRYPLI
	1					SELKQILLDTYYGRDYVEDRLS
		İ				FLLAVATLNGERLDGEMSEGK
						LVMPFRHVSDAFEQTRETIGVR
		İ				GNNAINDMVRQRLLN\RFTSEQ
						AEGNAIYRLTPLGIGITDYYIRQ
						REFSTLRLSMQLSIVAGELKRA
		1				ADAAEEGGDEFHWHRNVYAPL
						KYSVAEIFDSIDLTQRLMDEQQ
						QQVKDDIAQLLNKDWRAAISS CELLLSETSGTLRELQDTLEAA
						GDKLQANLLRIQDATMTHDDL
						HFVDRLVFDLQSKLDRIISWGQ
						QSIDLWIGYDRHVHKFIRTGID
					:	MDKKPRPSRRCL
24194	54562	A	24332	5	220	GLSWLNYC*ASTLTISLRCATR
						AVPLTRIRCRPRLLPSRRERTAL
						PCIYVKIAVTLLTATCASCVRR
						WIPA
24195	54563	Α	24333	3	180	
24196	54564	Α	24334	229	564	PGFKPRHSSWQRLPSLGHLPLY
						GSSVFTLLNLATAHSSGPCLLQ
						LELSFHSPSTTAVCRRHRPTADF
						HPSESGRVSHVLLIQ/QRCPLLL
24107	54565	 	24225	140	404	WIRLKACHCSCTAKCPGSSQLS
24197	54565	A	24335	213	404	 SCEHSVAHGRGIMDLSLLKALS
24198	54566	Α	24336	213	1306	EADAIASSEQEVRQIL\LEEADR
		1				LQKEVRVDGLGSVLIRLNESTG
						PKVMICAHMDEVGFMVRSISRE
						GAIDVLPVGNVRMAARQLQPV
						RITTREECKIPGLLDGDRQGND
						VSAMRVDIGARSYDEVMQAGI
						RPGDRVTFDTTFQVLPHQRVM
						GKAFDDRLGCYLLVTQLRELH
						DAELPAEVWLVASSSEEVGLR
						GGQTS\TRAVSPDVAIVLDTAC
					1	WAKNFDYGAANHRQIGNGPM
						LVLSDKSLIAPPKLTAWVETVA
						AEIGVPLQADMFSNGGTDGGA
						VHLTGTGVPTVVMGPATRHGH
						CAASIADCRDILHMQQLLSAFI
		1				HKSNELGGVLRLRIQICRDGNE
24199	54567	В	24337	1	1075	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24200	54568	Ā	24338	438	1462	QIVQCLDVPSYTFIWFAVLDPIY
						GDSVSTGLVVAIISIIVNAITIPIG
						LYLLNPSSGADGKKNSNLSALI
						SAAKEPVVWAPVLATILVLVG
						VKIPAAWDPTFNLIAKANSGVA
						VFAAGLTLAAHKFEFSAEIAYN
						TFLKLILMPLALLLVGMACHLN
					İ	SEHLQMI\TLASALPPAFSLKLIY
						CAA/ARLMCYTRTGTTSLAITLT
						ETVTQKLTGSRPRISSYGRNLM
						WPLAWYKDNGKPHLCSQLETT
						RIHSTAAQGRSSGSLRRGLGAS
						PFYRQHDQARSEGQIPITRIPMG
		1]	LAQRAPKGRAQPRGLTAPFTN
		1				GALGQPQGRGTSKQASQTRPR
		1				ASAWPHTPT
24201	54569	Α	24339	30	177	GSGYQRCHNNAASDRLRSYRK
		1				QPAER/WLAPPIDSITSPPASRRA
						LILLY
24202	54570	Α	24340	3350	3430	
24203	54571	Α	24341	425	824	SISGSENACFNAT\NHSGRYSQY
1		1				GACLTSSSCRKGQLARRGLSYQ
		1				SPHAPADPAFPAARRSPVFCLS
						DDGGHKSPRDMSPDKPSRHPD
						QVALTRRIPSPPAVALPSVVKR
	<u> </u>	<u> </u>			0.110	TVHEGQGDDDESNPSPLSPDVH
24204	54572	В	24342	11	2118	
24205 24206	54573 54574	A	24343 24344	730	1101	
24207	54575	B	24344	1	1992	
24208	54576	A	24346	880	1103	TMMRRLISRQPVLPLQKCFSVLI
27200	31370	``	2 13 10		1103	SNPWRVFGGTGGGNQER/CRY
						AEESVAETRLCP*CRLYAEYDP
	1					GSGCAGLSL
24209	54577	A	24347	11	952	MITLRKLPLAVAVAAGVMSAQ
						AMAVDFHGYARSGIGWTGSGG
						EQQCFQTTGAQSKYRLGNECE
	1	ŀ				TYAELKLGQEVWKEGDKSFYF
	1					DTNVAYSVAQQNDWEATDPAF
						REANVQGKNLIEWLPGSTIWA
						GKRFYORHDVHMIDFYYWDIS
						GPGAGLENIDVGFGKLSLAATR
						SSEAGGSSSFASNHIYDYS/NEN
				1		AND/VFDVRLARWDSPGGPLTG
						FPPSALACTSGVSGQKGVPGPP
						EAQVTFATPKKPGKEACSSIQV
						LNRGYRPPTFSYNTGVRRSARD
						YYPRRQRSISTNSPSRRWKGKL
						NSRRCSTGNRPLVGAVS
	<u> </u>		L	1	L	

Sequence	POSSIBLE NUCLEOTIDE TKLAMRVAIFDSV AASPEHSRFAVLW ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSCHTSTASLSVH LRFDHSSSRRRLCY DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GHMLRILDHGAIS
Sequence	TKLAMRVAIFDSV AASPEHSRFAVLW ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
24210 54578 A 24348 805 1274 ASFATTRPV	AASPEHSRFAVLW ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
VGQRYWGS. ILASIFR/DRI. MS*AWYWQ QFFSPARSAS. IALGYRSSES. RDYAHLHGI C 24211 54579 A 24349 275 490 24212 54580 A 24350 1 696 24213 54581 A 24351 I 735 MFTPDYRSV MNQPDVMP HMIDFYYWI GFGKLSLAA SNNIYDYTN QMEINPGGT RDNYRLGKG KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF QNIRTIYLAK	AASPEHSRFAVLW ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
ILASIFR/DRI. MS*AWYWQ QFFSPARSAS IALGYRSSES RDYAHLHGI C	ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
ILASIFR/DRI. MS*AWYWQ QFFSPARSAS IALGYRSSES RDYAHLHGI C	ASLRVQATVKS*K GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
MS*AWYWQ QFFSPARSAS IALGYRSSES RDYAHLHGI C	GCFRNWKSSPA*Y SSHFAGEPVSLHSP SSSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
QFFSPARSAS IALGYRSSES RDYAHLHGI C	SSHFAGEPVSLHSP SSSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
IALGYRSSES RDYAHLHGI C	SSCHTSTASLSVH LRFDHSSSRRRLCY SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
RDYAHLHGI C	SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
C	SRVLGSGPGYASR DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
24212 54580 A 24350 1 696	DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
24213 54581 A 24351 I 735 MFTPDYRSV MNQPDVMP HMIDFYYWI GFGKLSLAA SNNIYDYTN QMEINPGGT RDNYRLGKG KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF QNIRTIYLAK	DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
MNQPDVMP HMIDFYYWI GFGKLSLAA SNNIYDYTN QMEINPGGT RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF	DIPGKRFYQRHDV DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
HMIDFYYWI GFGKLSLAA SNNIYDYTN QMEINPGGT RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF	DISGPGAGLENIDV TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
GFGKLSLAA SNNIYDYTN QMEINPGGT RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W PIMNTGMKF QNIRTIYLAK	TRSSEAGGSSSFA ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
SNNIYDYTN QMEINPGGT RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF QNIRTIYLAK	ETANDVFDVRLA LELGVDYGRANL GLSQGSGVAFDNE
QMEINPGGT RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W' PIMNTGMKF QNIRTIYLAK	LELGVDYGRANL : GLSQGSGVAFDNE
RDNYRLGKO KFAYNINNN MGDNWDMI NDHG\TK\W PIMNTGMKF QNIRTIYLAK	GLSQGSGVAFDNE
KFAYNINNN MGDNWDMI NDHG\TK\W PIMNTGMKF QNIRTIYLAK	
MGDNWDMI NDHG\TK\W PIMNTGMKF QNIRTIYLAK	GHMLRILDHGAIS
NDHG\TK\W PIMNTGMKF QNIRTIYLAK	OIII.IDIGODIIO
PIMNTGMKF QNIRTIYLAK	MYVGMYQDIHWD
QNIRTIYLAK	WT\AGIRP\MFKWT
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	GYE/NVESQGPG
	P*QPDVMPDIPGK
	łMIDFYYWDISGP
GAGLENIDV GAGLENIDV	GFGKLSLAATRSS
EAGGSSSFA:	SNNIYDYTNETAN
DVFDVRLAC	MEINPGGTLELG
VDYGRANLI	RDNYRLGKGLSQG
SGVAFDNEK	FAYNINNNGHML
RILDHGAISM	IGDNWDMMYVG
MYQDIHWD	NDHGHQVGGPAG
	HHEHRDEIRYEKL
NPRARDKTF	VQFTSQNPDRC
24214 54582 A 24352 I 3414 MIVQPLAIIG	SHFDSIAAGNLAR
PIAVYGRNE	ITAIFASLKTMQQ
ALRGTVSDV	RKGSQEMHIGIAE
IVAGNNDLS	SRTEQQAASLAQT
AASMEQLTA	ATVGQNADNARQ
ASELAKNAA	TTAQAGGVQVST
MTHTMQEIA	TSSQKIGDIISVID
GIAFQTNILA	LNAAVEAARAGE
	AGEVRNLASRSAQ
	EESVNRVQQGSKL
	IDIVSSVTRVNDIM
GEIASASEEC	
24215 54583 A 24353 1 2493	